GenCore version 5.1.6

OM protein - protein search, using sw model

Run on: February 27, 2005, 20:02:49 ; Search time 115.869 Seconds
(without alignments)
1387.712 Million cell updates/sec

Title: US-10-040-647-6
Perfect score: 1728
Sequence: 1 MGARGALLLALLLARAGLRK.....PSWPLLFFPLLWALPLLGPV 314

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1612378 seqs, 512079187 residues

Total number of hits satisfying chosen parameters: 1612378

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : UniProt_03:*
1: uniprot_sprot:*
2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1723	99.7	314	1	TEST_HUMAN	Q9y6m0 homo sapien
2	1152	66.7	324	1	TEST_MOUSE	Q9jhj7 mus musculu
3	1152	66.7	336	2	Q80YD8	Q80yd8 mus musculu
4	1128.5	65.3	328	2	Q80Z40	Q80z40 rattus norv
5	881	51.0	318	2	Q7RTY9	Q7rty9 homo sapien
6	850	49.2	322	2	Q920S2	Q920s2 mus musculu
7	817	47.3	282	2	Q9D4I3	Q9d4i3 mus musculu
8	605	35.0	328	2	Q6BEA2	Q6bea2 rattus norv
9	597.5	34.6	321	2	Q6IE60	Q6ie60 rattus norv
10	596.5	34.5	321	1	TRYG_HUMAN	Q9nrr2 homo sapien
11	595.5	34.5	321	2	Q96RZ8	Q96rz8 homo sapien
12	586	33.9	719	2	Q6DJ90	Q6dj90 xenopus tro
13	585	33.9	328	2	Q8BJR6	Q8bjr6 mus musculu
14	579.5	33.5	343	1	PSS8_HUMAN	Q16651 homo sapien
15	575.5	33.3	389	2	Q9PVX7	Q9pvx7 xenopus lae
16	573	33.2	290	1	PR27_HUMAN	Q9bqr3 homo sapien
17	572.5	33.1	311	2	Q80XZ3	Q80xz3 rattus norv
18	570	33.0	330	2	Q6NVR7	Q6nvr7 xenopus tro
19	567.5	32.8	339	2	Q99L44	Q99l44 mus musculu
20	567.5	32.8	342	1	PSS8_RAT	Q9es87 rattus norv
21	566.5	32.8	331	2	Q8R1A6	Q8r1a6 mus musculu
22	566	32.8	344	2	Q640F8	Q640f8 xenopus lae
23	565	32.7	321	2	Q6GNK3	Q6gnk3 xenopus lae
24	563.5	32.6	331	2	Q80X17	Q80x17 mus musculu
25	563.5	32.6	342	1	PSS8_MOUSE	Q9esd1 mus musculu
26	561	32.5	317	2	Q9DGR3	Q9dgr3 xenopus lae
27	554.5	32.1	311	1	TRYG_MOUSE	Q9qul7 mus musculu
28	552	31.9	340	2	Q8BJV6	Q8bjv6 mus musculu
29	551	31.9	336	2	Q7RTY5	Q7rty5 homo sapien
30	546.5	31.6	312	2	Q7M755	Q7m755 mus musculu
31	542	31.4	799	2	Q6PF94	Q6pf94 mus musculu
32	542	31.4	811	1	TMS6_MOUSE	Q9dbi0 mus musculu
33	541.5	31.3	309	2	Q6DHH4	Q6dhh4 brachydanio
34	539.5	31.2	307	2	Q7TML0	Q7tml0 mus musculu
35	539	31.2	273	1	TRYT_SHEEP	Q9xsm2 ovis aries
36	538.5	31.2	297	2	O88781	O88781 rattus ratt
37	536.5	31.0	317	1	BSS4_HUMAN	Q9gzn4 homo sapien
38	536.5	31.0	334	2	Q6UXE0	Q6uxe0 homo sapien
39	527.5	30.5	310	1	DISP_MOUSE	Q9qyz9 mus musculu
40	525	30.4	306	1	BSS4_MOUSE	Q9er10 mus musculu
41	523.5	30.3	270	1	TRYT_MERUN	P50342 meriones un
42	517.5	29.9	471	2	Q8CFE0	Q8cfe0 mus musculu
43	517	29.9	273	2	Q9XSM1	Q9xsm1 ovis aries
44	517	29.9	275	1	TRYT_PIG	Q9n2d1 sus scrofa
45	513.5	29.7	638	1	KAL_MOUSE	P26262 mus musculu

ALIGNMENTS

RESULT 1
TEST_HUMAN
ID TEST_HUMAN STANDARD; PRT; 314 AA.
AC Q9Y6M0; Q9NS34; Q9P2V6;
DT 16-OCT-2001 (Rel. 40, Created)
DT 16-OCT-2001 (Rel. 40, Last sequence update)
DT 25-OCT-2004 (Rel. 45, Last annotation update)
DE Testisin precursor (EC 3.4.21.-) (Eosinophil serine protease 1) (ESP-
DE 1) (UNQ266/PRO303).
GN Name=PRSS21; Synonyms=ESP1, TEST1;
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A. (ISOFORM 1).
RC TISSUE=Eosinophil;
RX MEDLINE=99045401; PubMed=9826525; DOI=10.1006/bbrc.1998.9645;
RA Inoue M., Kanbe N., Kurosawa M., Kido H.;
RT "Cloning and tissue distribution of a novel serine protease esp-1 from
RT human eosinophils.";
RL Biochem. Biophys. Res. Commun. 252:307-312(1998).
RN [2]
RP SEQUENCE FROM N.A. (ISOFORM 3).
RX MEDLINE=20068805; PubMed=10600542; DOI=10.1006/bbrc.1999.1870;
RA Inoue M., Isobe M., Itoyama T., Kido H.;
RT "Structural analysis of esp-1 gene (PRSS 21).";
RL Biochem. Biophys. Res. Commun. 266:564-568(1999).
RN [3]
RP SEQUENCE FROM N.A. (ISOFORMS 1 AND 2).
RC TISSUE=Cervical carcinoma;
RX MEDLINE=99323395; PubMed=10397266;
RA Hooper J.D., Nicol D.L., Dickinson J.L., Eyre H.J., Scarman A.L.,
RA Normyle J.F., Stuttgen M.A., Douglas M.L., Loveland K.A.,
RA Sutherland G.R., Antalis T.M.;
RT "Testisin, a new human serine proteinase expressed by premeiotic
RT testicular germ cells and lost in testicular germ cell tumors.";
RL Cancer Res. 59:3199-3205(1999).
RN [4]
RP SEQUENCE FROM N.A.
RX MEDLINE=20461760; PubMed=11004480; DOI=10.1016/S0167-4781(00)00071-3;
RA Hooper J.D., Bowen N., Marshall H., Cullen L.M., Sood R., Daniels R.,
RA Stuttgen M.A., Normyle J.F., Higgs D.R., Kastner D.L., Ogbourne S.M.,
RA Pera M.F., Jazwinska E.C., Antalis T.M.;
RT "Localization, expression and genomic structure of the gene encoding
RT the human serine protease testisin.";
RL Biochim. Biophys. Acta 1492:63-71(2000).
RN [5]
RP SEQUENCE FROM N.A. (ISOFORM 1).
RX MEDLINE=22887296; PubMed=12975309; DOI=10.1101/gr.1293003;
RA Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J.,
RA Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P.,
RA Eaton D., Foster J., Grimaldi C., Gu Q., Hass P.E., Heldens S.,
RA Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J.,

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RA   Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J.,
RA   Seshagiri S., Simmons L., Singh J., Smith V., Stinson J., Vagts A.,
RA   Vandlen R., Watanabe C., Wieand D., Woods K., Xie M.-H., Yansura D.,
RA   Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I.,
RA   Godowski P., Gray A.;
RT   "The secreted protein discovery initiative (SPDI), a large-scale
RT   effort to identify novel human secreted and transmembrane proteins: a
RT   bioinformatics assessment.";
RL   Genome Res. 13:2265-2270(2003).
CC   -!- FUNCTION: Could regulate proteolytic events associated with
CC       testicular germ cell maturation.
CC   -!- SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor
CC       (Potential).
CC   -!- ALTERNATIVE PRODUCTS:
CC       Event=Alternative splicing; Named isoforms=3;
CC       Name=1; Synonyms=L;
CC         IsoId=Q9Y6M0-1; Sequence=Displayed;
CC       Name=2; Synonyms=S;
CC         IsoId=Q9Y6M0-2; Sequence=VSP_005389;
CC       Name=3;
CC         IsoId=Q9Y6M0-3; Sequence=VSP_005390;
CC   -!- TISSUE SPECIFICITY: Expressed predominantly in premeiotic
CC       testicular germ cells, mostly late pachytene and diplotene
CC       spermatocytes.
CC   -!- SIMILARITY: Belongs to the peptidase S1 family.
CC   ---------------------------------------------------------------------
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CC   or send an email to license@isb-sib.ch).
CC   ---------------------------------------------------------------------
DR   EMBL; AF058300; AAD41588.1; -.
DR   EMBL; AB031329; BAA83520.1; -.
DR   EMBL; AB031330; BAA83521.1; -.
DR   EMBL; AB031331; BAA89532.1; -.
DR   EMBL; AF058301; AAF79019.1; -.
DR   EMBL; AF058301; AAF79020.1; -.
DR   EMBL; AY359034; AAQ89393.1; -.
DR   HSSP; P00734; 1GHW.
DR   MEROPS; S01.011; -.
DR   Genew; HGNC:9485; PRSS21.
DR   MIM; 608159; -.
DR   GO; GO:0005737; C:cytoplasm; TAS.
DR   GO; GO:0005624; C:membrane fraction; TAS.
DR   GO; GO:0005886; C:plasma membrane; TAS.
DR   GO; GO:0008236; F:serine-type peptidase activity; TAS.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; 1.
DR   PROSITE; PS00135; TRYPSIN_SER; 1.
KW   Alternative splicing; Glycoprotein; GPI-anchor; Hydrolase;
KW   Lipoprotein; Serine protease; Signal; Zymogen.
FT   SIGNAL       1     19       Potential.
FT   PROPEP      20     41       Potential.
FT   CHAIN       42    288       Testisin.
FT   PROPEP     289    314       Removed in mature form (Potential).
FT   ACT_SITE    82     82       Charge relay system (Potential).
FT   ACT_SITE   137    137       Charge relay system (Potential).
FT   ACT_SITE   238    238       Charge relay system (Potential).
FT   DISULFID    33    157       Potential.
FT   DISULFID    67     83       Potential.
FT   DISULFID   171    244       Potential.
FT   DISULFID   204    223       Potential.
FT   DISULFID   234    262       Potential.
FT   LIPID      288    288       GPI-anchor amidated serine (Potential).
FT   CARBOHYD   167    167       N-linked (GlcNAc. . .) (Potential).
FT   CARBOHYD   200    200       N-linked (GlcNAc. . .) (Potential).
FT   CARBOHYD   273    273       N-linked (GlcNAc. . .) (Potential).
FT   VARSPLIC    87     88       Missing (in isoform 2).
FT                               /FTId=VSP_005389.
FT   VARSPLIC   222    235       Missing (in isoform 3).
FT                               /FTId=VSP_005390.
SQ   SEQUENCE   314 AA;  34884 MW;  E738CF73F6B56E98 CRC64;

  Query Match             99.7%;  Score 1723;  DB 1;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1.8e-145;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:|||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314


RESULT 2
TEST_MOUSE
ID   TEST_MOUSE     STANDARD;      PRT;   324 AA.
AC   Q9JHJ7; Q9DA14;
DT   16-OCT-2001 (Rel. 40, Created)
DT   16-OCT-2001 (Rel. 40, Last sequence update)
DT   25-OCT-2004 (Rel. 45, Last annotation update)
DE   Testisin precursor (EC 3.4.21.-) (Tryptase 4).
GN   Name=Prss21;
OS   Mus musculus (Mouse).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX   NCBI_TaxID=10090;
RN   [1]
RP   SEQUENCE FROM N.A.
RC   STRAIN=129/Sv;
RX   MEDLINE=21153229; PubMed=11231276;
RA   Scarman A.L., Hooper J.D., Boucaut K.J., Sit M.-L., Webb G.C.,
RA   Normyle J.F., Antalis T.M.;
RT   "Organization and chromosomal localization of the murine Testisin gene
RT   encoding a serine protease temporally expressed during
RT   spermatogenesis.";
RL   Eur. J. Biochem. 268:1250-1258(2001).
RN   [2]
RP   SEQUENCE FROM N.A.
RC   STRAIN=BALB/c; TISSUE=Testis;
RX   PubMed=11259427; DOI=10.1074/jbc.M010422200;
RA   Wong G.W., Li L., Madhusudhan M.S., Krilis S.A., Gurish M.F.,
RA   Rothenberg M.E., Sali A., Stevens R.L.;
RT   "Tryptase 4, a new member of the chromosome 17 family of mouse serine
RT   proteases.";
RL   J. Biol. Chem. 276:20648-20658(2001).
RN   [3]
RP   SEQUENCE OF 3-324 FROM N.A.
RC   STRAIN=C57BL/6J; TISSUE=Testis;
RX   MEDLINE=22354683; PubMed=12466851; DOI=10.1038/nature01266;
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RA   Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kondo S.,
RA   Nikaido I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H.,
RA   Yagi K., Tomaru Y., Hasegawa Y., Nogami A., Schonbach C., Gojobori T.,
RA   Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,
RA   Schriml L.M., Kanapin A., Matsuda H., Batalov S., Beisel K.W.,
RA   Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,
RA   Dalla E., Dragani T.A., Fletcher C.F., Forrest A., Frazer K.S.,
RA   Gaasterland T., Gariboldi M., Gissi C., Godzik A., Gough J.,
RA   Grimmond S., Gustincich S., Hirokawa N., Jackson I.J., Jarvis E.D.,
RA   Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,
RA   Konagaya A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,
RA   Maglott D.R., Maltais L., Marchionni L., McKenzie L., Miki H.,
RA   Nagashima T., Numata K., Okido T., Pavan W.J., Pertea G., Pesole G.,
RA   Petrovsky N., Pillai R., Pontius J.U., Qi D., Ramachandran S.,
RA   Ravasi T., Reed J.C., Reed D.J., Reid J., Ring B.Z., Ringwald M.,
RA   Sandelin A., Schneider C., Semple C.A., Setou M., Shimada K.,
RA   Sultana R., Takenaka Y., Taylor M.S., Teasdale R.D., Tomita M.,
RA   Verardo R., Wagner L., Wahlestedt C., Wang Y., Watanabe Y., Wells C.,
RA   Wilming L.G., Wynshaw-Boris A., Yanagisawa M., Yang I., Yang L.,
RA   Yuan Z., Zavolan M., Zhu Y., Zimmer A., Carninci P., Hayatsu N.,
RA   Hirozane-Kishikawa T., Konno H., Nakamura M., Sakazume N., Sato K.,
RA   Shiraki T., Waki K., Kawai J., Aizawa K., Arakawa T., Fukuda S.,
RA   Hara A., Hashizume W., Imotani K., Ishii Y., Itoh M., Kagawa I.,
RA   Miyazaki A., Sakai K., Sasaki D., Shibata K., Shinagawa A.,
RA   Yasunishi A., Yoshino M., Waterston R., Lander E.S., Rogers J.,
RA   Birney E., Hayashizaki Y.;
RT   "Analysis of the mouse transcriptome based on functional annotation of
RT   60,770 full-length cDNAs.";
RL   Nature 420:563-573(2002).
CC   -!- FUNCTION: Could regulate proteolytic events associated with
CC       testicular germ cell maturation.
CC   -!- SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor
CC       (Potential).
CC   -!- TISSUE SPECIFICITY: Testis.
CC   -!- DEVELOPMENTAL STAGE: Expressed in post-meiotic testicular germ
CC       cells.
CC   -!- SIMILARITY: Belongs to the peptidase S1 family.
CC   -!- CAUTION: Ref.3 sequence differs from that shown due to a stop
CC       codon in position 315.
CC   ---------------------------------------------------------------------
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CC   or send an email to license@isb-sib.ch).
CC   ---------------------------------------------------------------------
DR   EMBL; AF304012; AAK29360.1; -.
DR   EMBL; AY005145; AAG02255.1; -.
DR   EMBL; AF176209; AAF64407.2; -.
DR   EMBL; AF226710; AAF64428.2; -.
DR   EMBL; AK006271; -; NOT_ANNOTATED_CDS.
DR   HSSP; P00734; 1GHW.
DR   MEROPS; S01.011; -.
DR   MGD; MGI:1916698; Prss21.
DR   GO; GO:0005624; C:membrane fraction; IDA.
DR   GO; GO:0004252; F:serine-type endopeptidase activity; IDA.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; 1.
DR   PROSITE; PS00135; TRYPSIN_SER; 1.
KW   Glycoprotein; GPI-anchor; Hydrolase; Lipoprotein; Serine protease;
KW   Signal; Zymogen.
FT   SIGNAL       1     21       Potential.
FT   PROPEP      22     54       Potential.
FT   CHAIN       55    298       Testisin.
FT   PROPEP     299    324       Removed in mature form (Potential).
FT   ACT_SITE    95     95       Charge relay system (Potential).
FT   ACT_SITE   147    147       Charge relay system (Potential).
FT   ACT_SITE   248    248       Charge relay system (Potential).
FT   DISULFID    46    167       Potential.
FT   DISULFID    80     96       Potential.
FT   DISULFID   181    254       Potential.
FT   DISULFID   214    233       Potential.
FT   DISULFID   244    272       Potential.
FT   LIPID      298    298       GPI-anchor amidated asparagine
FT                               (Potential).
FT   CARBOHYD   170    170       N-linked (GlcNAc. . .) (Potential).
FT   CARBOHYD   177    177       N-linked (GlcNAc. . .) (Potential).
FT   CARBOHYD   210    210       N-linked (GlcNAc. . .) (Potential).
FT   CARBOHYD   283    283       N-linked (GlcNAc. . .) (Potential).
FT   CONFLICT   275    275       P -> H (in Ref. 3).
SQ   SEQUENCE   324 AA;  36175 MW;  56DC59E84F3C3CD4 CRC64;

  Query Match           66.7%;  Score 1152;  DB 1;  Length 324;
  Best Local Similarity  65.0%;  Pred. No. 1.9e-94;
  Matches  212;  Conservative   38;  Mismatches   60;  Indels   16;  Gaps    4;

Qy          1 MGARGALLLALL--LARAGL----------RKPESQEAAPLSGPCGRRVITSRIVGGED 47
              |||||  |: ||  :| | :            ||| ||   |||||| | | |||||:|
Db          1 MGARGKTLVPLLVVVATAAMALQSTYLQVDPEKPELQEPDLLSGPCGHRTIPSRIVGGDD 60

Qy         48 AELGRWPWQGSLRLWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMP 107
              |||||||||||||:| :|:|| :||: || |||||||:   | :||  | ||||:||| |
Db         61 AELGRWPWQGSLRVWGNHLCGATLLNRRWVLTAAHCFQ--KD-NDPFDWTVQFGELTSRP 117

Qy        108 SFWSLQAYYTRYFVSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFEN 167
              | |:|||| || : :|:|||:|    | ||||:|||:||||   ||||||  ||::|||
Db        118 SLWNLQAYSNRYQIEDIFLSPKYSEQYPNDIALLKLSSPVTYNNFIQPICLLNSTYKFEN 177

Qy        168 RTDCWVTGWGYIKEDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNA 227
              |||||||||| | |||:||||:||||||||||||||||:: |  || :|:|||||||
Db        178 RTDCWVTGWGAIGEDESLPSPNTLQEVQVAIINNSMCNHMYKKPDFRTNIWGDMVCAGTP 237

Qy        228 QGGKDACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQ 287
              :|||||||||||||::| :|||:||||||:||||||||||||||||: |||  | :
Db        238 EGGKDACFGDSGGPLACDQDTVWYQVGVVSWGIGCGRPNRPGVYTNISHHYNWIQSTMIR 297

Qy        288 SGMSQPDPSWPLLFFPLLWALPLLGP 313
              :|: :|||   |||  | ||  || |
Db        298 NGLLRPDPVPLLLFLTLAWASSLLRP 323


RESULT 3
Q80YD8
ID   Q80YD8      PRELIMINARY;      PRT;   336 AA.
AC   Q80YD8;
DT   01-JUN-2003 (TrEMBLrel. 24, Created)
DT   01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Prss21 protein (Fragment).
GN   Name=Prss21;
OS   Mus musculus (Mouse).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX   NCBI_TaxID=10090;
RN   [1]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Testis;
RX   MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA   Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA   Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA   Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA   Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA   Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA   Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA   Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA   Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA   Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA   Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA   Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
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RA   Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA   Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA   Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA   Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA   Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA   Jones S.J., Marra M.A.;
RT   "Generation and initial analysis of more than 15,000 full-length human
RT   and mouse cDNA sequences.";
RL   Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN   [2]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Testis;
RA   Strausberg R.;
RL   Submitted (MAR-2003) to the EMBL/GenBank/DDBJ databases.
CC   -!- SIMILARITY: Belongs to peptidase family S1.
DR   EMBL; BC049588; AAH49588.1; -.
DR   HSSP; P00766; 1CHG.
DR   MGD; MGI:1916698; Prss21.
DR   GO; GO:0005615; C:extracellular space; TAS.
DR   GO; GO:0005624; C:membrane fraction; IDA.
DR   GO; GO:0004252; F:serine-type endopeptidase activity; IDA.
DR   GO; GO:0007283; P:spermatogenesis; TAS.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   SMART; SM00020; Tryp_SPc; 1.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
DR   PROSITE; PS00135; TRYPSIN_SER; 1.
KW   Hydrolase; Protease; Serine protease.
FT   NON_TER       1       1
SQ   SEQUENCE   336 AA;  37361 MW;  E5206FEDBE55C670 CRC64;

  Query Match             66.7%;  Score 1152;  DB 2;  Length 336;
  Best Local Similarity   65.0%;  Pred. No. 1.9e-94;
  Matches  212;  Conservative   38;  Mismatches   60;  Indels   16;  Gaps    4;

Qy          1 MGARGALLLALL--LARAGL----------RKPESQEAAPLSGPCGRRVITSRIVGGED 47
              |||||  |: ||  :| | :           ||| ||   |||||| | | ||||||:|
Db         13 MGARGKTLVPLLVVVATAAMALQSTYLQVDPEKPELQEPDLLSGPCGHRTIPSRIVGGDD 72

Qy         48 AELGRWPWQGSLRLWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMP 107
              |||||||||||||:| :|:|| :||: || |||||||:   | :||  | ||||:||| |
Db         73 AELGRWPWQGSLRVWGNHLCGATLLNRRWVLTAAHCFQ--KD-NDPFDWTVQFGELTSRP 129

Qy        108 SFWSLQAYYTRYFVSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFEN 167
              | |:||||  || : :|:|||:|    | ||||:|||:||||  ||||||  ||::|||
Db        130 SLWNLQAYSNRYQIEDIFLSPKYSEQYPNDIALLKLSSPVTYNNFIQPICLLNSTYKFEN 189

Qy        168 RTDCWVTGWGYIKEDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNA 227
              |||||||||| | |||:||||:||||||||||||||||:: |  || :|:|||||||
Db        190 RTDCWVTGWGAIGEDESLPSPNTLQEVQVAIINNSMCNHMYKKPDFRTNIWGDMVCAGTP 249

Qy        228 QGGKDACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQ 287
              :||||||||||||||::| :|||:||||||:||||||||||||||||: |||  | :
Db        250 EGGKDACFGDSGGPLACDQDTVWYQVGVVSWGIGCGRPNRPGVYTNISHHYNWIQSTMIR 309

Qy        288 SGMSQPDPSWPLLFFPLLWALPLLGP 313
              :|: :|||   |||  | ||  || |
Db        310 NGLLRPDPVPLLLFLTLAWASSLLRP 335


RESULT 4
Q80Z40
ID   Q80Z40      PRELIMINARY;      PRT;   328 AA.
AC   Q80Z40;
DT   01-JUN-2003 (TrEMBLrel. 24, Created)
DT   01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Eosinophil serine protease-1.
GN   Name=resp-1;
OS   Rattus norvegicus (Rat).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
OX   NCBI_TaxID=10116;
RN   [1]
RP   SEQUENCE FROM N.A.
RC   STRAIN=Sprague-Dawley; TISSUE=Testis;
RA   Nakamura Y., Inoue M., Okumura Y., Shiota M., Nishikawa M., Arase S.,
RA   Kido H.;
RT   "Cloning, expression analysis, and tissue distribution of esp-
RT   1/testisin, a membrane-type serine protease from the rat.";
RL   J. Med. Invest. 50:78-86(2003).
CC   -!- SIMILARITY: Belongs to peptidase family S1.
DR   EMBL; AB074516; BAC57949.1; -.
DR   HSSP; P00766; 1CHG.
DR   MEROPS; S01.011; -.
DR   GO; GO:0004263; F:chymotrypsin activity; IEA.
DR   GO; GO:0008233; F:peptidase activity; IEA.
DR   GO; GO:0004295; F:trypsin activity; IEA.
DR   GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   SMART; SM00020; Tryp_SPc; 1.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
DR   PROSITE; PS00135; TRYPSIN_SER; 1.
KW   Hydrolase; Protease; Serine protease.
SQ   SEQUENCE   328 AA;  36631 MW;  7F6C0B204802B963 CRC64;

  Query Match             65.3%;  Score 1128.5;  DB 2;  Length 328;
  Best Local Similarity   63.5%;  Pred. No. 2.4e-92;
  Matches  205;  Conservative   40;  Mismatches   59;  Indels   19;  Gaps    3;

Qy          1 MGARGALLLALLLARAGL----------------RKPESQEAAPLSGPCGRRVITSRIVG 44
              | |||  |: ||:    :                 ||| |||  |||||| | | ||||||
Db          1 MSARGKTLVPLLVVVVAVEVTLQSTSSHVKPVDPEKPELQEANLLSGPCGHRTIPSRIVG 60

Qy         45 GEDAELGRWPWQGSLRLWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLT 104
              ||:|||||||||||:| :|:|| :||: || |||||||:    | :||  | ||||:||
Db         61 GEEAELGRWPWQGSLRVWGNHLCGATLLNRRWVLTAAHCFQ--KD-NDPFDWTVQFGELT 117

Qy        105 SMPSFWSLQAYYTRYFVSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFE 164
              | || |:|||| || : :|:|||:|    |:||||:|||:||||:  ||||||  ||::
Db        118 SRPSLWNLQAYSNRYQIEDIFLSPKYTEQFPHDIALLKLSSPVTYSNFIQPICLLNSTYK 177

Qy        165 FENRTDCWVTGWGYIKEDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCA 224
              | |||||||||| | |||:|| |: |||||||||:||||| |  || :|:||||||
Db        178 FANRTDCWVTGWGAIGEDESLPLPNNLQEVQVAIINNTMCNHLFKKPDFRINIWGDMVCA 237

Qy        225 GNAQGGKDACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKL 284
              |: :|||||||||||||| ||:| :|||:||||||:|||||||||||||||||: ||:
Db        238 GSPEGGKDACFGDSGGPLVCNQDTVWYQVGVVSWGIGCGRPNRPGVYTNISHHYNWIRLT 297

Qy        285 MAQSGMSQPDPSWPLLFFPLLWA 307
              | ::|| :|||:  |||  | ||
Db        298 MIRNGMLRPDPAPLLLFLTLAWA 320


RESULT 5
Q7RTY9
ID   Q7RTY9      PRELIMINARY;      PRT;   318 AA.
AC   Q7RTY9;
DT   01-MAR-2004 (TrEMBLrel. 26, Created)
DT   01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Testis serine protease 1 precursor.
GN   Name=TESSP1;
OS   Homo sapiens (Human).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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OX   NCBI_TaxID=9606;
RN   [1]
RP   SEQUENCE FROM N.A.
RX   MEDLINE=22722134; PubMed=12838346;
RA   Puente X.S., Sanchez L.M., Overall C.M., Lopez-Otin C.;
RT   "Human and mouse proteases: a comparative genomic approach.";
RL   Nat. Rev. Genet. 4:544-558(2003).
CC   -!- SIMILARITY: Belongs to peptidase family S1.
CC   -!- MISCELLANEOUS: The sequence shown here is derived from an
CC       EMBL/GenBank/DDBJ third party annotation (TPA) entry.
DR   EMBL; BN000124; CAD67575.1; -.
DR   HSSP; P08709; 1FAK.
DR   MEROPS; S01.327; -.
DR   GO; GO:0004263; F:chymotrypsin activity; IEA.
DR   GO; GO:0008233; F:peptidase activity; IEA.
DR   GO; GO:0004295; F:trypsin activity; IEA.
DR   GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
DR   PROSITE; PS00135; TRYPSIN_SER; 1.
KW   Hydrolase; Protease; Serine protease; Signal.
FT   SIGNAL        1     20       Potential.
FT   CHAIN        21    318       testis serine protease 1.
SQ   SEQUENCE   318 AA;  35078 MW;  D792213630CF15C3 CRC64;

  Query Match             51.0%;  Score 881;  DB 2;  Length 318;
  Best Local Similarity   53.7%;  Pred. No. 2.9e-70;
  Matches  173;  Conservative   45;  Mismatches   84;  Indels   20;  Gaps    5;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGP--------------CGRRVITSRIVGGE 46
              |||||||||||||||||| ||  :  |  :||              || | | : : ||
Db          1 MGARGALLLALLLARAGLGKP-GELGALQAGPGAARRPGGGGREEACGHREIHALVAGGV 59

Qy         47 DAELGRWPWQGSLRLWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSM 106
              ::  |||||| ||||    | || |||| || |:|||||: :     || | || |:|||
Db         60 ESARGRWPWQASLRLRRRHRCGGSLLSRRWVLSAAHCFQKH---YYPSEWTVQLGELTSR 116

Qy        107 PSFWSLQAYYTRYFVSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFE 166
              |: |:|:|| :|| | :| ::|   ||    ||||::|::|||  :|||||:::||| |
Db        117 PTPWNLRAYSSRYKVQDIIVNPDALGVLRNDIALLRLASSVTYNAYIQPICIESSTFNFV 176

Qy        167 NRTDCWVTGWGYIK-EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAG 225
              :| |||||||| |       || |: |:| || |:||: ||:|| : | |  |:  | :|||
Db        177 HRPDCWVTGWGLISPSGTPLPPPYNLREAQVTILNNTRCNYLFEQPSSRSMIWDSMFCAG 236

Qy        226 NAQGGKDACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLM 285
                |  | | ||||||| |:|||||||:|:||||: ||:||||||||||| :| ||:::|
Db        237 AEDGSVDTCKGDSGGPLVCDKDGLWYQVGIVSWGMDCGQPNRPGVYTNISVYFHWIRRVM 296

Qy        286 AQSGMSQPDPSWPLLFFPLLWA 307
              : |   :|:||  ||   ||||
Db        297 SHS-TPRPNPSQLLLLLALLWA 317


RESULT 6
Q920S2
ID   Q920S2      PRELIMINARY;      PRT;   322 AA.
AC   Q920S2;
DT   01-DEC-2001 (TrEMBLrel. 19, Created)
DT   01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Testis serine protease-1.
GN   Name=4931440B09Rik; Synonyms=tessp-1;
OS   Mus musculus (Mouse).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX   NCBI_TaxID=10090;
RN   [1]
RP   SEQUENCE FROM N.A.
RA   Matsui H., Takano N., Takahashi T.;
RL   Submitted (SEP-2000) to the EMBL/GenBank/DDBJ databases.
CC   -!- SIMILARITY: Belongs to peptidase family S1.
DR   EMBL; AB049453; BAB68561.1; -.
DR   HSSP; P00766; 1CHG.
DR   MEROPS; S01.417; -.
DR   MGD; MGI:1918253; 4931440B09Rik.
DR   GO; GO:0005615; C:extracellular space; TAS.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   SMART; SM00020; Tryp_SPc; 1.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
DR   PROSITE; PS00135; TRYPSIN_SER; 1.
KW   Hydrolase; Protease; Serine protease.
SQ   SEQUENCE   322 AA;  36218 MW;  ED55F9A199A3E491 CRC64;

  Query Match             49.2%;  Score 850;  DB 2;  Length 322;
  Best Local Similarity   52.8%;  Pred. No. 1.7e-67;
  Matches  172;  Conservative   43;  Mismatches   93;  Indels   18;  Gaps    6;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAP----------LSGPCGRRVIT-SRIVGGEDA 48
              || :| :|| |||    | || |:| :           || |||||  | |||||| ::
Db          1 MGIQGPVLLLLLLC-VMLGKPGSREESQAADLKSTDIKLLSMPCGRRNDTRSRIVGGIES 59

Qy         49 ELGRWPWQGSLRLWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPS 108
              ||||| ||||  || || |||| || |||||||  |    ||  | || ||||| ||
Db         60 MQGRWPWQASLRLKKSHRCGGSLLSRRWVLTAAHCFRKY---LDPEKWTVQLGQLTSKPS 116

Qy        109 FWSLQAYYTRYFVSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENR 168
              :|: :||  || | :| ::      | :|:||::|:: ||| | |||:|:| |||  :::
Db        117 YWNRKAYSGRYRVKDIIVNSEDKLKS-HDLALLRLASSVTYNKDIQPVCVQPSTFTSQHQ 175

Qy        169 TDCWVTGWGYIKED-EALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNA 227
              ||||||| ::|| : || |: |:||||:|:||| |  ||  :|     |  |: |||
Db        176 PRCWVTGWGVLQEDLKPLPPPYHLREVQVSILNNSRCQELFEIFSLHHLITKDVFCAGAE 235

Qy        228 QGGKDACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQ 287
              |  | | ||||||| || |||||||:||||:|||||| ||:|||:||:: ||: :|
Db        236 DGSADTCSGDSGGPLVCNMDGLWYQIGIVSWGIGCGRPNLPGIYTNVSHYYNWIETMMIL 295

Qy        288 SGMSQPDPSWPLLFFPLLWALPLLGP 313
              :|  : | : |||   || |  || |
Db        296 NGAVRRDLALPLLSITLLQAPWLLRP 321


RESULT 7
Q9D4I3
ID   Q9D4I3      PRELIMINARY;      PRT;   282 AA.
AC   Q9D4I3;
DT   01-JUN-2001 (TrEMBLrel. 17, Created)
DT   01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Mus musculus adult male testis cDNA, RIKEN full-length enriched
DE   library, clone:4931440B09 product:TESTIS SERINE PROTEASE-1
DE   homolog.
GN   Name=4931440B09Rik;
OS   Mus musculus (Mouse).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX   NCBI_TaxID=10090;
RN   [1]
RP   SEQUENCE FROM N.A.
RC   STRAIN=C57BL/6J; TISSUE=Testis;
RX   MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
RA   Carninci P., Hayashizaki Y.;
RT   "High-efficiency full-length cDNA cloning.";
RL   Meth. Enzymol. 303:19-44(1999).
RN   [2]
RP   SEQUENCE FROM N.A.
RC   STRAIN=C57BL/6J; TISSUE=Testis;
RX   MEDLINE=21085660; PubMed=11217851; DOI=10.1038/35055500;
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RA   RIKEN FANTOM Consortium;
RT   "Functional annotation of a full-length mouse cDNA collection.";
RL   Nature 409:685-690(2001).
RN   [3]
RP   SEQUENCE FROM N.A.
RC   STRAIN=C57BL/6J; TISSUE=Testis;
RA   The FANTOM Consortium,
RA   the RIKEN Genome Exploration Research Group Phase I & II Team;
RT   "Analysis of the mouse transcriptome based on functional annotation of
RT   60,770 full-length cDNAs.";
RL   Nature 420:563-573(2002).
RN   [4]
RP   SEQUENCE FROM N.A.
RC   STRAIN=C57BL/6J; TISSUE=Testis;
RX   MEDLINE=20499374; PubMed=11042159; DOI=10.1101/gr.145100;
RA   Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata K., Itoh M.,
RA   Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.;
RT   "Normalization and subtraction of cap-trapper-selected cDNAs to
RT   prepare full-length cDNA libraries for rapid discovery of new genes.";
RL   Genome Res. 10:1617-1630(2000).
RN   [5]
RP   SEQUENCE FROM N.A.
RC   STRAIN=C57BL/6J; TISSUE=Testis;
RX   MEDLINE=20530913; PubMed=11076861; DOI=10.1101/gr.152600;
RA   Shibata K., Itoh M., Aizawa K., Nagaoka S., Sasaki N., Carninci P.,
RA   Konno H., Akiyama J., Nishi K., Kitsunai T., Tashiro H., Itoh M.,
RA   Sumi N., Ishii Y., Nakamura S., Hazama M., Nishine T., Harada A.,
RA   Yamamoto R., Matsumoto H., Sakaguchi S., Ikegami T., Kashiwagi K.,
RA   Fujiwake S., Inoue K., Togawa Y., Izawa M., Ohara E., Watahiki M.,
RA   Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J.,
RA   Okazaki Y., Muramatsu M., Inoue Y., Kira A., Hayashizaki Y.;
RT   "RIKEN integrated sequence analysis (RISA) system-384-format
RT   sequencing pipeline with 384 multicapillary sequencer.";
RL   Genome Res. 10:1757-1771(2000).
RN   [6]
RP   SEQUENCE FROM N.A.
RC   STRAIN=C57BL/6J; TISSUE=Testis;
RA   Adachi J., Aizawa K., Akahira S., Akimura T., Arai A., Aono H.,
RA   Arakawa T., Bono H., Carninci P., Fukuda S., Fukunishi Y., Furuno M.,
RA   Hanagaki T., Hara A., Hayatsu N., Hiramoto K., Hiraoka T., Hori F.,
RA   Imotani K., Ishii Y., Itoh M., Izawa M., Kasukawa T., Kato H.,
RA   Kawai J., Kojima Y., Konno H., Kouda M., Koya S., Kurihara C.,
RA   Matsuyama T., Miyazaki A., Nishi K., Nomura K., Numazaki R., Ohno M.,
RA   Okazaki Y., Okido T., Owa C., Saito H., Saito R., Sakai C., Sakai K.,
RA   Sano H., Sasaki D., Shibata K., Shibata Y., Shinagawa A., Shiraki T.,
RA   Sogabe Y., Suzuki H., Tagami M., Tagawa A., Takahashi F., Tanaka T.,
RA   Tejima Y., Toya T., Yamamura T., Yasunishi A., Yoshida K., Yoshino M.,
RA   Muramatsu M., Hayashizaki Y.;
RL   Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.
CC   -!- SIMILARITY: Belongs to peptidase family S1.
DR   EMBL; AK016509; BAB30277.1; -.
DR   HSSP; P00766; 1CHG.
DR   MEROPS; S01.417; -.
DR   MGD; MGI:1918253; 4931440B09Rik.
DR   GO; GO:0005615; C:extracellular space; TAS.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   SMART; SM00020; Tryp_SPc; 1.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
DR   PROSITE; PS00135; TRYPSIN_SER; 1.
KW   Hydrolase; Protease; Serine protease.
SQ   SEQUENCE   282 AA;  31997 MW;  48D89B74F750BAB4 CRC64;

  Query Match           47.3%;  Score 817;  DB 2;  Length 282;
  Best Local Similarity   55.3%;  Pred. No. 1.3e-64;
  Matches  157;  Conservative   39;  Mismatches   82;  Indels    6;  Gaps    4;

Qy         32 PCGRRVIT-SRIVGGEDAELGRWPWQGSLRLWDSHVCGVSLLSHRWALTAAHCFETYSDL 90
              |||||  | |||||| ::  |||||| ||||  || || |||| || |||||||  |

Db          2 PCGRRNDTRSRIVGGIESMQGRWPWQASLRLKKSHRCGGSLLSRRWVLTAAHCFRKY--- 58

Qy         91 SDPSGWMVQFGQLTSMPSFWSLQAYYTRYFVSNIYLSPRYLGNSPYDIALVKLSAPVTYT 150
              ||  | || ||||| ||:|: :||  || | :| ::       | :|:||::|:: |||

Db         59 LDPEKWTVQLGQLTSKPSYWNRKAYSGRYRVKDIIVNSEDKLKS-HDLALLRLASSVTYN 117

Qy        151 KHIQPICLQASTFEFENRTDCWVTGWGYIKED-EALPSPHTLQEVQVAIINNSMCNHLFL 209
              | |||:|:| |||  :::  ||||||| ::|| : || |: |:||||:|:||| |  ||

Db        118 KDIQPVCVQPSTFTSQHQPRCWVTGWGVLQEDLKPLPPPYHLREVQVSILNNSRCQELFE 177

Qy        210 KYSFRKDIFGDMVCAGNAQGGKDACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPG 269
               :|    |  |: |||    |  | | ||||||| || |||||||:||||:|||||| ||

Db        178 IFSLHHLITKDVFCAGAEDGSADTCSGDSGGPLVCNMDGLWYQIGIVSWGIGCGRPNLPG 237

Qy        270 VYTNISHHFEWIQKLMAQSGMSQPDPSWPLLFFPLLWALPLLGP 313
              :|||:||:: ||: :|  :|  : | : |||   || |  || |
Db        238 IYTNVSHYYNWIETMMILNGAVRRDLALPLLSITLLQAPWLLRP 281


RESULT 8
Q6BEA2
ID   Q6BEA2      PRELIMINARY;      PRT;   328 AA.
AC   Q6BEA2;
DT   25-OCT-2004 (TrEMBLrel. 28, Created)
DT   25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
DT   25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
DE   Marapsin.
OS   Rattus norvegicus (Rat).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
OX   NCBI_TaxID=10116;
RN   [1]
RP   SEQUENCE FROM N.A.
RC   STRAIN=Wister; TISSUE=Limbal epithelium;
RA   Adachi W., Norman B., Davis J., Piatigorsky J.;
RT   "Comprehensive analysis of gene expression of rat corneal epithelium
RT   and limbal epithelium.";
RL   Submitted (FEB-2003) to the EMBL/GenBank/DDBJ databases.
CC   -!- SIMILARITY: Belongs to peptidase family S1.
DR   EMBL; AB101677; BAC81507.1; -.
DR   GO; GO:0004263; F:chymotrypsin activity; IEA.
DR   GO; GO:0004295; F:trypsin activity; IEA.
DR   GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   SMART; SM00020; Tryp_SPc; 1.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
DR   PROSITE; PS00135; TRYPSIN_SER; 1.
KW   Hydrolase; Protease; Serine protease.
SQ   SEQUENCE   328 AA;  35803 MW;  9235EECAF62C1358 CRC64;

  Query Match           35.0%;  Score 605;  DB 2;  Length 328;
  Best Local Similarity   43.6%;  Pred. No. 1.3e-45;
  Matches  137;  Conservative   40;  Mismatches   99;  Indels   38;  Gaps   10;

Qy          6 ALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRLWDSH 65
              ||||  || |:|    |: |      ||   : :|:|||||  | |||| |::   :|
Db          8 ALLLLPLLLRSGTEGAEAMRA------CGHPRMFNRMVGGEDALEGEWPWQVSIQRNGAH 61

Qy         66 VCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQ-----AYYTRYF 120
              || ||::  | |||||||   |: || | : |  | |       ||     | |
Db         62 FCGGSLIAPTWVLTAAHCF---SNTSDISIYQVLLGAL-------KLQQPGPHALYVP-- 109

Qy        121 VSNIYLSPRYLG-NSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYI 179
              |  :   | | |  |  |:|||:|  |||:||:| |:||   :  |::  :|||||||
Db        110 VKRVKSHPEYQGMASSADVALVELQVPVTFTKYILPVCLPDPSVVFKSGMNCWVTGWGSP 169

Qy        180 KEDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFR----KDIFGDMVCAGNAQGGKDACF 235
               | : ||:|  ||:: | :|:   || |: | :      | |  ||:||| |:| ||||
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Db        170 SEQDRLPNPRILQKLAVPLIDTPKCNLLYSKDAEADIQLKTIKDDMLCAGFAEGKKDACK 229

Qy        236 GDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLM--------AQ 287
              ||||||| |   |  | | ||:||| || | ||||||  :: |::|| :::        |
Db        230 GDSGGPLVCLVDQSWVQAGVISWGEGCARRNRPGVYIRVASHYQWIHQIIPELQFQGRAG 289

Qy        288 SGMSQPDP-SW-PL 299
              |   | ||  | ||
Db        290 SQQQQRDPRGWQPL 303


RESULT 9
Q6IE60
ID   Q6IE60      PRELIMINARY;      PRT;   321 AA.
AC   Q6IE60;
DT   05-JUL-2004 (TrEMBLrel. 27, Created)
DT   05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
DT   05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE   Marapsin precursor.
GN   Name=mpn;
OS   Rattus norvegicus (Rat).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
OX   NCBI_TaxID=10116;
RN   [1]
RP   SEQUENCE FROM N.A.
RC   STRAIN=Sprague-Dawley;
RX   PubMed=15060002; DOI=10.1101/gr.1946304;
RA   Puente X.S., Lopez-Otin C.;
RT   "A genomic analysis of rat proteases and protease inhibitors.";
RL   Genome Res. 14:609-622(2004).
CC   -!- SIMILARITY: Belongs to peptidase family S1.
CC   -!- MISCELLANEOUS: The sequence shown here is derived from an
CC       EMBL/GenBank/DDBJ third party annotation (TPA) entry.
DR   EMBL; BN000333; CAE48388.1; -.
DR   HSSP; P00734; 1B7X.
DR   GO; GO:0004263; F:chymotrypsin activity; IEA.
DR   GO; GO:0008233; F:peptidase activity; IEA.
DR   GO; GO:0004295; F:trypsin activity; IEA.
DR   GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   SMART; SM00020; Tryp_SPc; 1.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
DR   PROSITE; PS00135; TRYPSIN_SER; 1.
KW   Hydrolase; Protease; Serine protease; Signal.
FT   SIGNAL        1     26       Potential.
FT   CHAIN        27    321       marapsin.
SQ   SEQUENCE   321 AA;  35053 MW;  A87735DF7A52F9E1 CRC64;

  Query Match             34.6%;  Score 597.5;  DB 2;  Length 321;
  Best Local Similarity   43.3%;  Pred. No. 6.1e-45;
  Matches  136;  Conservative   38;  Mismatches   95;  Indels   45;  Gaps   10;

Qy          6 ALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRLWDSH 65
              ||||  || |:||               ||   : :|:|||||  | |||| |::   :|
Db          8 ALLLLPLLLRSGL------------PACGHPRMFNRMVGGEDALEGEWPWQVSIQRNGAH 55

Qy         66 VCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQ-----AYYTRYF 120
               || ||::  | ||||||| |     || | : |  | |          ||      | |
Db         56 FCGGSLIAPTWVLTAAHCFST----SDISIYQVLLGAL-------KLQQPGPHALYVP-- 102

Qy        121 VSNIYLSPRYLG-NSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYI 179
              |  :   | | |   |:|||:|  |||:||:| |:||   :  |::  :|||||||
Db        103 VKRVKSHPEYQGMASSADVALVELQVPVTFTKYILPVCLPDPSVVFKSGMNCWVTGWGSP 162

Qy        180 KEDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFR----KDIFGDMVCAGNAQGGKDACF 235
              | : ||:|  ||:: | :|:    || |: | :       | |  ||:||| |:| ||||
Db        163 SEQDRLPNPRILQKLAVPLIDTPKCNLLYSKDAEADIQLKTIKDDMLCAGFAEGKKDACK 222

Qy        236 GDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLM--------AQ 287
              ||||||| |   |  | | ||:||| || | ||||||  :: |::|| :::        |
Db        223 GDSGGPLVCLVDQSWVQAGVISWGEGCARRNRPGVYIRVASHYQWIHQIIPELQFQGRAG 282

Qy        288 SGMSQPDP-SW-PL 299
              |   | ||  | ||
Db        283 SQQQQRDPRGWQPL 296


RESULT 10
TRYG_HUMAN
ID   TRYG_HUMAN     STANDARD;      PRT;   321 AA.
AC   Q9NRR2; Q9C015; Q9NRQ8; Q9UBB2;
DT   16-OCT-2001 (Rel. 40, Created)
DT   16-OCT-2001 (Rel. 40, Last sequence update)
DT   25-OCT-2004 (Rel. 45, Last annotation update)
DE   Tryptase gamma precursor (EC 3.4.21.-) (Transmembrane tryptase).
GN   Name=TPSG1; Synonyms=TMT;
OS   Homo sapiens (Human).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX   NCBI_TaxID=9606;
RN   [1]
RP   SEQUENCE FROM N.A. (VARIANTS GAMMA-1 AND GAMMA-2).
RX   MEDLINE=20302813; PubMed=10843716;
RA   Caughey G.H., Raymond W.W., Blount J.L., Hau L.W., Pallaoro M.,
RA   Wolters P.J., Verghese G.M.;
RT   "Characterization of human gamma-tryptases, novel members of the
RT   chromosome 16p mast cell tryptase and prostasin gene families.";
RL   J. Immunol. 164:6566-6575(2000).
RN   [2]
RP   SEQUENCE FROM N.A.
RX   MEDLINE=99452974; PubMed=10521469; DOI=10.1074/jbc.274.43.30784;
RA   Wong G.W., Tang Y., Feyfant E., Sali A., Li L., Li Y., Huang C.,
RA   Friend D.S., Krilis S.A., Stevens R.L.;
RT   "Identification of a new member of the tryptase family of mouse and
RT   human mast cell proteases which possesses a novel COOH-terminal
RT   hydrophobic extension.";
RL   J. Biol. Chem. 274:30784-30793(1999).
RN   [3]
RP   SEQUENCE OF 220-321 FROM N.A.
RA   Mittman S., Agnew W.S.;
RT   "Organization and alternative splicing of CACNA1H.";
RL   Submitted (JAN-2001) to the EMBL/GenBank/DDBJ databases.
CC   -!- SUBCELLULAR LOCATION: Membrane-anchored (Potential).
CC   -!- TISSUE SPECIFICITY: Expressed in many tissues.
CC   -!- POLYMORPHISM: There are two alleles; gamma-I and gamma-II which
CC       differ by 5 residues.
CC   -!- SIMILARITY: Belongs to the peptidase S1 family. Tryptase
CC       subfamily.
CC   ---------------------------------------------------------------------
CC   This SWISS-PROT entry is copyright. It is produced through a collaboration
CC   between the Swiss Institute of Bioinformatics and the EMBL outstation -
CC   the European Bioinformatics Institute. There are no restrictions on its
CC   use by non-profit institutions as long as its content is in no way
CC   modified and this statement is not removed. Usage by and for commercial
CC   entities requires a license agreement (See http://www.isb-sib.ch/announce/
CC   or send an email to license@isb-sib.ch).
CC   ---------------------------------------------------------------------
DR   EMBL; AF191031; AAF76457.1; -.
DR   EMBL; AF195508; AAF76458.1; -.
DR   EMBL; AF175759; AAF03697.1; -.
DR   EMBL; AF175522; AAF03695.1; -.
DR   EMBL; AF223563; AAG48852.2; -.
DR   HSSP; P00760; 1EZX.
DR   MEROPS; S01.028; -.
DR   Genew; HGNC:14134; TPSG1.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
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DR   SMART; SM00020; Tryp_SPc; 1.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; 1.
DR   PROSITE; PS00135; TRYPSIN_SER; FALSE_NEG.
KW   Glycoprotein; Hydrolase; Polymorphism; Serine protease; Signal;
KW   Transmembrane; Zymogen.
FT   SIGNAL        1     19       Potential.
FT   CHAIN        20    321       Tryptase gamma.
FT   CHAIN        20     36       Tryptase gamma light chain.
FT   CHAIN        38    321       Tryptase gamma heavy chain.
FT   TRANSMEM    284    304       Potential.
FT   ACT_SITE     78     78       Charge relay system (By similarity).
FT   ACT_SITE    125    125       Charge relay system (By similarity).
FT   ACT_SITE    222    222       Charge relay system (By similarity).
FT   DISULFID     26    145       Interchain (Potential).
FT   DISULFID     63     79       By similarity.
FT   DISULFID    159    228       By similarity.
FT   DISULFID    192    210       By similarity.
FT   DISULFID    218    246       By similarity.
FT   CARBOHYD     85     85       N-linked (GlcNAc. . .) (Potential).
FT   VARIANT      60     60       M -> V (in gamma-II).
FT                                /FTId=VAR_012097.
FT   VARIANT     126    126       I -> M (in gamma-II).
FT                                /FTId=VAR_012098.
FT   VARIANT     132    132       S -> T (in gamma-II).
FT                                /FTId=VAR_012099.
FT   VARIANT     204    204       L -> I (in gamma-II).
FT                                /FTId=VAR_012100.
FT   VARIANT     288    288       L -> F (in gamma-II).
FT                                /FTId=VAR_012101.
FT   CONFLICT    160    160       W -> S (in Ref. 1).
SQ   SEQUENCE   321 AA;  33827 MW;  FFF7B06E3C4A962D CRC64;

  Query Match             34.5%;  Score 596.5;  DB 1;  Length 321;
  Best Local Similarity   42.7%;  Pred. No. 7.5e-45;
  Matches  138;  Conservative   40;  Mismatches  100;  Indels   45;  Gaps   10;

Qy          3 ARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVIT---SRIVGGEDAELGRWPWQGSL 59
              | ||  | ||||  |:          |   |||  ::    |||||  |  | |||| ||
Db          2 ALGACGLLLLLAVPGV------SLRTLQPGCGRPQVSDAGGRIVGGHAAPAGAWPWQASL 55

Qy         60 RLWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQL--TSMPSFWSLQAYYT 117
              ||   |||| |||| :| |||||||    : ||   : |  |:|  |  | | :::
Db         56 RLRRMHVCGGSLLSPQWVLTAAHCFSGSLNSSD---YQVHLGELEITLSPHFSTVRQII- 111

Qy        118 RYFVSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWG 177
                 :: ||     :  |||||:|| ||| :  | |:||  :: :|     |||||||
Db        112 ------LHSSPSGQPGTSGDIALVELSVPVTLSSRILPVCLPEASDDFCPGIRCWVTGWG 165

Qy        178 YIKEDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFG--------DMVCAGNAQG 229
              | :| | || |::|:||:|::::   |          |:|  |         ||:|   |:|
Db        166 YTREGEPLPPPYSLREVKVSVVDTETC---------RRDYPGPGGSILQPDMLC---ARG 213

Qy        230 GKDACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSG 289
               |||  |||||| |  :| | | |:|||| ||||||||||| :  :  ||:: : :||
Db        214 PGDACQDDSGGPLVCQVNGAWVQAGIVSWGEGCGRPNRPGVYTRVPAYVNWIRRHITASG 273

Qy        290 MSQPD-PSWPL---LFFPLLWAL 308
               |:   |  ||   || | |: |
Db        274 GSESGYPRLPLLAGLFLPGLFLL 296


RESULT 11
Q96RZ8
ID   Q96RZ8      PRELIMINARY;      PRT;   321 AA.
AC   Q96RZ8;
DT   01-DEC-2001 (TrEMBLrel. 19, Created)
DT   01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   HS transmembrane tryptase; gene name TMT; AF175522_1.
GN   Name=trpA;
OS   Homo sapiens (Human).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX   NCBI_TaxID=9606;
RN   [1]
RP   SEQUENCE FROM N.A.
RX   MEDLINE=21096910; PubMed=11157797; DOI=10.1093/hmg/10.4.339;
RA   Daniels R.J., Peden J.F., Lloyd C., Horsley S.W., Clark K.,
RA   Tufarelli C., Kearney L., Buckle V.J., Doggett N.A., Flint J.,
RA   Higgs D.R.;
RT   "Sequence, structure and pathology of the fully annotated terminal 2
RT   Mb of the short arm of human chromosome 16.";
RL   Hum. Mol. Genet. 10:339-352(2001).
CC   -!- SIMILARITY: Belongs to peptidase family S1.
DR   EMBL; AE006466; AAK61269.1; -.
DR   HSSP; P00760; 1EZX.
DR   GO; GO:0016021; C:integral to membrane; IEA.
DR   GO; GO:0004263; F:chymotrypsin activity; IEA.
DR   GO; GO:0008233; F:peptidase activity; IEA.
DR   GO; GO:0004295; F:trypsin activity; IEA.
DR   GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   SMART; SM00020; Tryp_SPc; 1.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
KW   Hydrolase; Protease; Serine protease; Transmembrane.
SQ   SEQUENCE   321 AA;  33829 MW;  FFF5089EDC4FC73D CRC64;

  Query Match             34.5%;  Score 595.5;  DB 2;  Length 321;
  Best Local Similarity   42.7%;  Pred. No. 9.2e-45;
  Matches  138;  Conservative   40;  Mismatches  100;  Indels   45;  Gaps   10;

Qy          3 ARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVIT---SRIVGGEDAELGRWPWQGSL 59
              | ||  | ||||  |:          |   |||  ::    |||||  |  | |||| ||
Db          2 ALGACGLLLLLAVPGV------SLRTLQPGCGRPQVSDAGGRIVGGHAAPAGAWPWQASL 55

Qy         60 RLWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQL--TSMPSFWSLQAYYT 117
              ||   |||| |||| :| |||||||    : ||   : |  |:|  |  | | :::
Db         56 RLRRVHVCGGSLLSPQWVLTAAHCFSGSLNSSD---YQVHLGELEITLSPHFSTVRQII- 111

Qy        118 RYFVSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWG 177
                 :: ||     :  |||||:|| ||| :  | |:||  :: :|     |||||||
Db        112 ------LHSSPSGQPGTSGDIALVELSVPVTLSSRILPVCLPEASDDFCPGIRCWVTGWG 165

Qy        178 YIKEDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFG--------DMVCAGNAQG 229
              | :| | || |::|:||:|::::   |          |:|  |         ||:|   |:|
Db        166 YTREGEPLPPPYSLREVKVSVVDTETC---------RRDYPGPGGSILQPDMLC---ARG 213

Qy        230 GKDACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSG 289
               |||  |||||| |  :| | | |:|||| ||||||||||| :  :  ||:: : :||
Db        214 PGDACQDDSGGPLVCQVNGAWVQAGIVSWGEGCGRPNRPGVYTRVPAYVNWIRRHITASG 273

Qy        290 MSQPD-PSWPLL---FFPLLWAL 308
               |:   |  |||   | | |: |
Db        274 GSESGYPRLPLLAGFFLPGLFLL 296


RESULT 12
Q6DJ90
ID   Q6DJ90      PRELIMINARY;      PRT;   719 AA.
AC   Q6DJ90;
DT   25-OCT-2004 (TrEMBLrel. 28, Created)
DT   25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
DT   25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
DE   Tmprss9-prov protein.
GN   Name=tmprss9-prov;
OS   Xenopus tropicalis (Western clawed frog) (Silurana tropicalis).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
OC   Xenopodinae; Xenopus.
OX   NCBI_TaxID=8364;
```

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RN   [1]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Whole body;
RX   MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA   Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA   Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA   Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA   Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA   Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA   Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA   Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA   Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA   Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA   Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA   Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA   Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA   Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA   Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA   Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA   Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA   Jones S.J., Marra M.A.;
RT   "Generation and initial analysis of more than 15,000 full-length human
RT   and mouse cDNA sequences.";
RL   Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN   [2]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Whole body;
RA   Klein S., Gerhard D.S.;
RL   Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.
CC   -!- SIMILARITY: Belongs to peptidase family S1.
DR   EMBL; BC075293; AAH75293.1; -.
DR   GO; GO:0004263; F:chymotrypsin activity; IEA.
DR   GO; GO:0004295; F:trypsin activity; IEA.
DR   GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   Pfam; PF00089; Trypsin; 2.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   SMART; SM00020; Tryp_SPc; 2.
DR   PROSITE; PS50240; TRYPSIN_DOM; 2.
DR   PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_2.
DR   PROSITE; PS00135; TRYPSIN_SER; 2.
KW   Hydrolase; Protease; Serine protease.
SQ   SEQUENCE   719 AA;  76098 MW;  906346A21D8834C6 CRC64;

  Query Match             33.9%;  Score 586;  DB 2;  Length 719;
  Best Local Similarity   43.3%;  Pred. No. 1.6e-43;
  Matches  117;  Conservative   43;  Mismatches   90;  Indels   20;  Gaps    7;

Qy         21 PESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRLWDSHVCGVSLLSHRWALTA 80
              |       |     ||  :::|||||| ||  | |||| |||    ||:|| |::  :| |||
Db        364 PTILSTTPAPPACGSPLVSSRIVGGTDAREGAWPWQVSLRYRGSHICGGSVIGTQWILTA 423

Qy         81 AHCFETYSDLSDPSGWMVQFGQ---LTSMPSFWSLQAYYT--RYPVSNIYLSPRYLGNSP 135
              |||||   :   || : |: |      : |:    :   ||  |  |:: : |    |
Db        424 AHCFE---NSQFPSDYEVRLGTYRLAQTSPN----EITYTVDRIIVNSQFDSSTLFG--- 473

Qy        136 YDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIKEDEALPSPHTLQEVQ 195
               ||||::|::|:||||:| |:|| :::  | :  :|||||| |      || | |||||
Db        474 -DIALIRLTSPITYTKYILPVCLPSTSNSFTDGMECWVTGWGTISLYVNLPYPKTLQEVM 532

Qy        196 VAIINNSMCN---HLFLKYSFRKDIF-GDMVCAGNAQGGKDACFGDSGGPLACNKDGLWY 251
                :|| : |:   |:    |   :|   | :|:| : ||||:| ||||||| |   |:||
Db        533 TPLINRTRCDQMYHIDSPVSASSEIIPSDQICSGYSAGGKDSCKGDSGGPLVCKLQGIWY 592

Qy        252 QIGVVSWGVGCGRPNRPGVYTNISHHFEWI 281
              |||:|||| ||    |||||| :  :: |:
Db        593 QIGIVSWGEGCAIAKRPGVYTLVPAYYSWV 622


RESULT 13
Q8BJR6
ID   Q8BJR6      PRELIMINARY;      PRT;   328 AA.
AC   Q8BJR6;
DT   01-MAR-2003 (TrEMBLrel. 23, Created)
DT   01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
DT   05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE   Mus musculus 3 days neonate thymus cDNA, RIKEN full-length enriched
DE   library, clone:A630023F11 product:similar to MARAPSIN (EC 3.4.21.-)
DE   (Pancreasin) (Channel-activating protease 2-like protein).
GN   Name=Mpn;
OS   Mus musculus (Mouse).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX   NCBI_TaxID=10090;
RN   [1]
RP   SEQUENCE FROM N.A.
RC   STRAIN=C57BL/6J; TISSUE=Thymus;
RX   MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
RA   Carninci P., Hayashizaki Y.;
RT   "High-efficiency full-length cDNA cloning.";
RL   Meth. Enzymol. 303:19-44(1999).
RN   [2]
RP   SEQUENCE FROM N.A.
RC   STRAIN=C57BL/6J; TISSUE=Thymus;
RX   MEDLINE=21085660; PubMed=11217851; DOI=10.1038/35055500;
RA   RIKEN FANTOM Consortium;
RT   "Functional annotation of a full-length mouse cDNA collection.";
RL   Nature 409:685-690(2001).
RN   [3]
RP   SEQUENCE FROM N.A.
RC   STRAIN=C57BL/6J; TISSUE=Thymus;
RA   The FANTOM Consortium,
RA   the RIKEN Genome Exploration Research Group Phase I & II Team;
RT   "Analysis of the mouse transcriptome based on functional annotation of
RT   60,770 full-length cDNAs.";
RL   Nature 420:563-573(2002).
RN   [4]
RP   SEQUENCE FROM N.A.
RC   STRAIN=C57BL/6J; TISSUE=Thymus;
RX   MEDLINE=20499374; PubMed=11042159; DOI=10.1101/gr.145100;
RA   Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata K., Itoh M.,
RA   Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.;
RT   "Normalization and subtraction of cap-trapper-selected cDNAs to
RT   prepare full-length cDNA libraries for rapid discovery of new genes.";
RL   Genome Res. 10:1617-1630(2000).
RN   [5]
RP   SEQUENCE FROM N.A.
RC   STRAIN=C57BL/6J; TISSUE=Thymus;
RX   MEDLINE=20530913; PubMed=11076861; DOI=10.1101/gr.152600;
RA   Shibata K., Itoh M., Aizawa K., Nagaoka S., Sasaki N., Carninci P.,
RA   Konno H., Akiyama J., Nishi K., Kitsunai T., Tashiro H., Itoh M.,
RA   Sumi N., Ishii Y., Nakamura S., Hazama M., Nishine T., Harada A.,
RA   Yamamoto R., Matsumoto H., Sakaguchi S., Ikegami T., Kashiwagi K.,
RA   Fujiwake S., Inoue K., Togawa Y., Izawa M., Ohara E., Watahiki M.,
RA   Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J.,
RA   Okazaki Y., Muramatsu M., Inoue Y., Kira A., Hayashizaki Y.;
RT   "RIKEN integrated sequence analysis (RISA) system-384-format
RT   sequencing pipeline with 384 multicapillary sequencer.";
RL   Genome Res. 10:1757-1771(2000).
RN   [6]
RP   SEQUENCE FROM N.A.
RC   STRAIN=C57BL/6J; TISSUE=Thymus;
RA   Adachi J., Aizawa K., Akimura T., Arakawa T., Bono H., Carninci P.,
RA   Fukuda S., Furuno M., Hanagaki T., Hara A., Hashizume W.,
RA   Hayashida K., Hayatsu N., Hiramoto K., Hiraoka T., Hirozane T.,
RA   Hori F., Imotani K., Ishii Y., Itoh M., Kagawa I., Kasukawa T.,
RA   Katoh H., Kawai J., Kojima Y., Kondo S., Konno H., Kouda M., Koya S.,
RA   Kurihara C., Matsuyama T., Miyazaki A., Murata M., Nakamura M.,
RA   Nishi K., Nomura K., Numazaki R., Ohno M., Ohsato N., Okazaki Y.,
RA   Saito R., Saitoh H., Sakai C., Sakai K., Sakazume N., Sano H.,
RA   Sasaki D., Shibata K., Shinagawa A., Shiraki T., Sogabe Y., Tagami M.,
RA   Tagawa A., Takahashi F., Takaku-Akahira S., Takeda Y., Tanaka T.,
RA   Tomaru A., Toya T., Yasunishi A., Muramatsu M., Hayashizaki Y.;
RL   Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases.
```

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RN   [7]
RP   SEQUENCE FROM N.A.
RC   STRAIN=C57BL/6; TISSUE=Bladder;
RX   MEDLINE=22439795; PubMed=12441343; DOI=10.1074/jbc.M209353200;
RA   Bhagwandin V.J., Hau L.W., Mallen-St  Clair J., Wolters P.J.,
RA   Caughey G.H.;
RT   "Structure and activity of human pancreasin, a novel tryptic serine
RT   peptidase expressed primarily by the pancreas.";
RL   J. Biol. Chem. 5:3363-3371(2003).
RN   [8]
RP   SEQUENCE FROM N.A.
RC   STRAIN=129/SvJ;
RA   Brathwaite M., Waeltz P., Schlessinger D., Nagaraja R.;
RL   Submitted (OCT-2002) to the EMBL/GenBank/DDBJ databases.
CC   -!- SIMILARITY: Belongs to peptidase family S1.
DR   EMBL; AK080281; BAC37864.1; -.
DR   EMBL; AF542056; AAO27572.1; -.
DR   EMBL; AY162410; AAO17162.1; -.
DR   HSSP; P00734; 1UVS.
DR   MEROPS; S01.074; -.
DR   MGD; MGI:2450123; Mpn.
DR   GO; GO:0004263; F:chymotrypsin activity; IEA.
DR   GO; GO:0008233; F:peptidase activity; IEA.
DR   GO; GO:0004295; F:trypsin activity; IEA.
DR   GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   SMART; SM00020; Tryp_SPc; 1.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
DR   PROSITE; PS00135; TRYPSIN_SER; 1.
KW   Hydrolase; Protease; Serine protease.
SQ   SEQUENCE   328 AA;  35789 MW;  DC0B20F1AB3EB840 CRC64;

  Query Match             33.9%;  Score 585;  DB 2;  Length 328;
  Best Local Similarity   44.2%;  Pred. No. 8.1e-44;
  Matches  129;  Conservative   41;  Mismatches   94;  Indels   28;  Gaps    8;

Qy          6 ALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRLWDSH 65
              ||||  || |:|       | |      ||   : :|:||||:|  | |||| |::   |
Db          8 ALLLLPLLLRSG------TEGARTLRACGHPKMFNRMVGGENALEGEWPWQVSIQRNGIH 61

Qy         66 VCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQ-----AYYTRYF 120
               || ||::  | |||||||   |: || | : |  | |         ||     | |
Db         62 FCGGSLIAPTWVLTAAHCF---SNTSDISIYQVLLGAL-------KLQQPGPHALYVP-- 109

Qy        121 VSNIYLSPRYLG-NSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYI 179
              |  :  :|:| |  |  |:|||:|  |||:| :|  |:||   :  ||:  :|||||||
Db        110 VKQVKSNPQYQGMASSADVALVELQGPVTFTNYILPVCLPDPSVIFESGMNCWVTGWGSP 169

Qy        180 KEDEALPSPHTLQEVQVAIINNSMCNHLF---LKYSFR-KDIFGDMVCAGNAQGGKDACF 235
              | : ||:|  ||:: | ||:   || |:   ::  |: | |  ||:||| |:| |||| 
Db        170 SEQDRLPNPRVLQKLAVPIIDTPKCNLLYNKDVESDFQLKTIKDDMLCAGFAEGKKDACK 229

Qy        236 GDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQ 287
              |||||| |  |  | | ||:||| || | ||||||  :: | :|| ::: :
Db        230 GDSGGPLVCLVDQSWVQAGVISWGEGCARRNRPGVYIRVTSHHKWIHQIIPE 281


RESULT 14
PSS8_HUMAN
ID   PSS8_HUMAN     STANDARD;      PRT;   343 AA.
AC   Q16651; Q9UCA3;
DT   01-NOV-1997 (Rel. 35, Created)
DT   01-NOV-1997 (Rel. 35, Last sequence update)
DT   25-OCT-2004 (Rel. 45, Last annotation update)
DE   Prostasin precursor (EC 3.4.21.-).
GN   Name=PRSS8;
OS   Homo sapiens (Human).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX   NCBI_TaxID=9606;
RN   [1]
RP   SEQUENCE FROM N.A., AND PARTIAL SEQUENCE.
RC   TISSUE=Prostate;
RX   MEDLINE=95286644; PubMed=7768952;
RA   Yu J.X., Chao L., Chao J.;
RT   "Molecular cloning, tissue-specific expression, and cellular
RT   localization of human prostasin mRNA.";
RL   J. Biol. Chem. 270:13483-13489(1995).
RN   [2]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Placenta;
RX   MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA   Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA   Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA   Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA   Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA   Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA   Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA   Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA   Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA   Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA   Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA   Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA   Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA   Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA   Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA   Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA   Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
RA   Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
RT   "Generation and initial analysis of more than 15,000 full-length human
RT   and mouse cDNA sequences.";
RL   Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN   [3]
RP   SEQUENCE OF 45-64.
RC   TISSUE=Semen;
RX   MEDLINE=94308140; PubMed=8034638;
RA   Yu J.X., Chao L., Chao J.;
RT   "Prostasin is a novel human serine proteinase from seminal fluid.
RT   Purification, tissue distribution, and localization in prostate
RT   gland.";
RL   J. Biol. Chem. 269:18843-18848(1994).
CC   -!- FUNCTION: Possesses a trypsin-like cleavage specificity.
CC   -!- SUBUNIT: Heterodimer of two chains, light and heavy, held by a
CC       disulfide bond.
CC   -!- SUBCELLULAR LOCATION: Membrane-bound. Secreted after cleavage of
CC       its C-terminus.
CC   -!- TISSUE SPECIFICITY: Found in prostate, liver, salivary gland,
CC       kidney, lung, pancreas, colon, bronchus and renal proximal tubular
CC       cells. In the prostate gland it may be synthesized in epithelial
CC       cells, secreted into the ducts, and excreted into the seminal
CC       fluid.
CC   -!- SIMILARITY: Belongs to the peptidase S1 family.
CC   ---------------------------------------------------------------------
CC   This SWISS-PROT entry is copyright. It is produced through a collaboration
CC   between the Swiss Institute of Bioinformatics and the EMBL outstation -
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CC   use by non-profit institutions as long as its content is in no way
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CC   or send an email to license@isb-sib.ch).
CC   ---------------------------------------------------------------------
DR   EMBL; L41351; AAC41759.1; -.
DR   EMBL; U33446; AAB19071.1; -.
DR   EMBL; BC001462; AAH01462.1; -.
DR   PIR; A57014; A57014.
DR   HSSP; P00760; 1EZX.
DR   MEROPS; S01.159; -.
DR   Genew; HGNC:9491; PRSS8.
DR   H-InvDB; HIX0012982; -.
DR   MIM; 600823; -.
DR   GO; GO:0005615; C:extracellular space; TAS.
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DR   GO; GO:0005886; C:plasma membrane; TAS.
DR   GO; GO:0008236; F:serine-type peptidase activity; TAS.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   SMART; SM00020; Tryp_SPc; 1.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; 1.
DR   PROSITE; PS00135; TRYPSIN_SER; 1.
KW   Direct protein sequencing; Glycoprotein; Hydrolase; Serine protease;
KW   Signal; Transmembrane; Zymogen.
FT   SIGNAL       1     29       Potential.
FT   PROPEP      30     32       Activation peptide.
FT   CHAIN       33     44       Prostasin light chain.
FT   CHAIN       45    322       Prostasin heavy chain.
FT   PROPEP     323    343
FT   TRANSMEM   320    340       Potential.
FT   DOMAIN      45    286       Serine protease.
FT   DISULFID    37    154       Interchain (By similarity).
FT   DISULFID    70     86       By similarity.
FT   DISULFID   168    244       By similarity.
FT   DISULFID   201    223       By similarity.
FT   DISULFID   234    262       By similarity.
FT   ACT_SITE    85     85       Charge relay system.
FT   ACT_SITE   134    134       Charge relay system.
FT   ACT_SITE   238    238       Charge relay system.
FT   CARBOHYD   159    159       N-linked (GlcNAc. . .) (Potential).
SQ   SEQUENCE   343 AA;  36431 MW;  98DD6447F5A8C1B2 CRC64;

  Query Match             33.5%;  Score 579.5;  DB 1;  Length 343;
  Best Local Similarity   41.0%;  Pred. No. 2.7e-43;
  Matches  141;  Conservative   45;  Mismatches  101;  Indels   57;  Gaps   12;

Qy          6 ALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRLWDSH 65
              |:|| | | |:|    |  ||     |||     :|| ||  |  |:|||| |:    |
Db         16 AILLYLGLLRSG-TGAEGAEA-----PCG-VAPQARITGGSSAVAGQWPWQVSITYEGVH 68

Qy         66 VCGVSLLSHRWALTAAHCFETYSDLSDPS-----GWMVQFG--QLTSMPSFWSLQAYYTR 118
              ||| ||:| :| |:|||||        ||      : |: |  || |    :|  |  :
Db         69 VCGGSLVSEQWVLSAAHCF--------PSEHHKEAYEVKLGAHQLDS----YSEDAKVST 116

Qy        119 YFVSNIYLSPRYL-GNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWG 177
                : :|   | ||   |  ||||::|| |:|::::|:|||| |:   | |   | |||||
Db        117 --LKDIIPHPSYLQEGSQGDIALLQLSRPITFSRYIRPICLPAANASFPNGLHCTVTGWG 174

Qy        178 YIKEDEALPSPHTLQEVQVAIINNSMCNHLF---LKYSFRKDIFGDMVCAGNAQGGKDAC 234
              ::    :| :|  ||:::| :|:   || |:    |      :  ||||||  :||||||
Db        175 HVAPSVSLLTPKPLQQLEVPLISRETCNCLYNIDAKPEEPHFVQEDMVCAGYVEGGKDAC 234

Qy        235 FGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQ--------KLMA 286
              |||||||:|  :|||| |:|||| ||  |||||||  | :  |||         :::
Db        235 QGDSGGPLSCPVEGLWYLTGIVSWGDACGARNRPGVYTLASSYASWIQSKVTELQPRVVP 294

Qy        287 QSGMSQPDPSW----------------PLLFFPLLWALPLLGP 313
              |:  |||| :                  |:|| ||  || || |
Db        295 QTQESQPDSNLCGSHLAFSSAPAQGLLRPILFLPLGLALGLLSP 338


RESULT 15
Q9PVX7
ID   Q9PVX7      PRELIMINARY;      PRT;   389 AA.
AC   Q9PVX7;
DT   01-MAY-2000 (TrEMBLrel. 13, Created)
DT   01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Epidermis specific serine protease.
GN   Name=Xepsin;
OS   Xenopus laevis (African clawed frog).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
OC   Xenopodinae; Xenopus.
OX   NCBI_TaxID=8355;
RN   [1]
RP   SEQUENCE FROM N.A.
RA   Yamada K.;
RL   Submitted (OCT-1998) to the EMBL/GenBank/DDBJ databases.
CC   -!- SIMILARITY: Belongs to peptidase family S1.
DR   EMBL; AB018694; BAA84941.1; -.
DR   HSSP; P00760; 1EZX.
DR   GO; GO:0004263; F:chymotrypsin activity; IEA.
DR   GO; GO:0008233; F:peptidase activity; IEA.
DR   GO; GO:0004295; F:trypsin activity; IEA.
DR   GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
DR   InterPro; IPR001254; Peptidase_S1.
DR   InterPro; IPR001314; Peptidase_S1A.
DR   InterPro; IPR009003; Pept_Ser_Cys.
DR   Pfam; PF00089; Trypsin; 1.
DR   PRINTS; PR00722; CHYMOTRYPSIN.
DR   SMART; SM00020; Tryp_SPc; 1.
DR   PROSITE; PS50240; TRYPSIN_DOM; 1.
DR   PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
DR   PROSITE; PS00135; TRYPSIN_SER; 1.
KW   Hydrolase; Protease; Serine protease.
SQ   SEQUENCE   389 AA;  42375 MW;  B31FB4A2F5D1F6E3 CRC64;

  Query Match             33.3%;  Score 575.5;  DB 2;  Length 389;
  Best Local Similarity   39.9%;  Pred. No. 6.9e-43;
  Matches  120;  Conservative   44;  Mismatches   84;  Indels   53;  Gaps    7;

Qy         33 CGRRVITSRIVGGEDAELGRWPWQGSLRLWDSHVCGVSLLSHRWALTAAHCFETYS---- 88
              ||  ||::||||| |:: | |||| ||      :|| |||:  | :|||||  ::
Db         17 CGVPVISNRIVGGMDSKRGEWPWQISLSYKSDSICGGSLLTDSWVMTAAHCIDSLDVSYY 76

Qy         89 -------DLSDPSGWMVQFG--QLTSMPSFWSLQAYYTRYFVSNIYLSPRYLGNSPYDIA 139
                    || |   | |   :|  | |                    :| |:|  |||
Db         77 TVYLGAYQLSAPDNSTVSRGVKSITKHPDF------------------QYEGSSG-DIA 116

Qy        140 LVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIKEDEALPSPHTLQEVQVAII 199
              |::|  |||:| :| |||| :   :|   | |||||||  |:|   | || |:|: :||||
Db        117 LIELEKPVTFTPYILPICLPSQDVQFAAGTMCWVTGWGNIQEGTPLISPKTIQKAEVAII 176

Qy        200 NNSMCNHLFLK-------YSFRKDIFGDMVCAGNAQGGKDACFGDSGGPLACNKDGLWYQ 252
              ::|:|  ::          :|| ::   ||||||  :|  ||| |||||||  || : :| |
Db        177 DSSVCGTMYESSLGYIPDFSFIQE---DMVCAGYKEGRIDACQGDSGGPLVCNVNNVWLQ 233

Qy        253 IGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLLFFPLLWALPLLG 312
              :|:|||| ||  |||||||  : :: :|::  :      |:  ||             | :|
Db        234 LGIVSWGYGCAEPNRPGVYTKVQYYQDWLKTNVPLIVFSEEGPS----------VAPSIG 283

Qy        313 P 313
              |
Db        284 P 284


Search completed: February 27, 2005, 20:18:32
Job time : 116.869 secs
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GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: March 4, 2005, 20:50:35 ; Search time 5114.55 Seconds
(without alignments)
10421.388 Million cell updates/sec

Title: US-10-040-647-5
Perfect score: 1100
Sequence: 1 CGCGGGAGAGGAGGCCATGG.....AAAAAAAAAAAAAAAAAAAA 1100

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 4708233 seqs, 24227607955 residues

Total number of hits satisfying chosen parameters: 9416466

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : GenEmbl:*
1: gb_ba:*
2: gb_htg:*
3: gb_in:*
4: gb_om:*
5: gb_ov:*
6: gb_pat:*
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8: gb_pl:*
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10: gb_ro:*
11: gb_sts:*
12: gb_sy:*
13: gb_un:*
14: gb_vi:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	DB	ID	Description
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5	1090.6	99.1	1100	6	BD173395	BD173395 Secreted
6	1090.6	99.1	1100	6	BD175429	BD175429 Secretory
7	1090.6	99.1	1100	6	AR410808	AR410808 Sequence
8	1090.6	99.1	1100	6	AR439172	AR439172 Sequence
9	1090.6	99.1	1100	6	AR473192	AR473192 Sequence
10	1090.6	99.1	1100	6	AR527178	AR527178 Sequence
11	1090.6	99.1	1100	6	AR566211	AR566211 Sequence
12	1090.6	99.1	1100	6	AX697665	AX697665 Sequence
13	1090.6	99.1	1100	6	BD075578	BD075578 Secretory
14	1090.6	99.1	1100	9	AY359034	AY359034 Homo sapi
15	1078	98.0	1094	6	AR253955	AR253955 Sequence
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19	1075.6	97.8	1082	9	AB031329	AB031329 Homo sapi
20	1071.8	97.4	1077	6	CQ714823	CQ714823 Sequence
21	1071.8	97.4	1077	6	AX370460	AX370460 Sequence
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25	1064.2	96.7	1085	6	E21853	E21853 Novel acido
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ALIGNMENTS

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LOCUS AR253956 1100 bp DNA linear PAT 20-DEC-2002
DEFINITION Sequence 5 from patent US 6479274.
ACCESSION AR253956
VERSION AR253956.1 GI:27302451
KEYWORDS .
SOURCE Unknown.
ORGANISM Unknown.
Unclassified.
REFERENCE 1 (bases 1 to 1100)
AUTHORS Antalis,T.M. and Hooper,J.D.
TITLE DNA molecules encoding human HELA2 or testisin serine proteinases
JOURNAL Patent: US 6479274-A 5 12-NOV-2002;
FEATURES Location/Qualifiers
source 1. .1100
/organism="unknown"
/mol_type="genomic DNA"
ORIGIN

Query Match 100.0%; Score 1100; DB 6; Length 1100;
Best Local Similarity 100.0%; Pred. No. 6.3e-267;
Matches 1100; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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LOCUS       BD172438               1100 bp    DNA     linear   PAT 18-FEB-2003
DEFINITION  Secreted and transmembrane polypeptides and nucleic acids encoding
            the same.
ACCESSION   BD172438
VERSION     BD172438.1  GI:28413738
KEYWORDS    JP 2002223786-A/211.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 1100)
  AUTHORS   Wood,W.I., Gurney,A.L., Goddard,A., Pennica,D., Zheng,J. and
            Yuan,J.
  TITLE     Secreted and transmembrane polypeptides and nucleic acids encoding
            the same
  JOURNAL   Patent: JP 2002223786-A 211 13-AUG-2002;
            GENENTECH INC
COMMENT     OS   Homo sapiens (human)
            PN   JP 2002223786-A/211
            PD   13-AUG-2002
            PF   18-DEC-2001 JP 2001385135
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            24-NOV-1997 US    60/066453,25-NOV-1997 US    60/066840 PI
            WILLIAM I WOOD,AUSTIN L GURNEY,AUDREY GODDARD,DIANE PENNICA, PI
            JIAN ZHENG,
            PI   JEAN YUAN
            PC   C12N15/09,C07K14/47,C07K16/18,C07K19/00,C12N1/19,C12N1/21, PC
             C12N5/10,
            PC
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Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
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RESULT 3
BD172757
LOCUS       BD172757               1100 bp    DNA     linear   PAT 18-FEB-2003
DEFINITION  Secreted and transmembrane polypeptides and nucleic acids encoding
            the same.
ACCESSION   BD172757
VERSION     BD172757.1  GI:28414061
KEYWORDS    JP 2002238586-A/211.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 1100)
  AUTHORS   Wood,W.I., Gurney,A.L., Goddard,A., Pennica,D., Zheng,J. and
            Yuan,J.
  TITLE     Secreted and transmembrane polypeptides and nucleic acids encoding
            the same
  JOURNAL   Patent: JP 2002238586-A 211 27-AUG-2002;
            GENENTECH INC
COMMENT     OS   Homo sapiens (human)
            PN   JP 2002238586-A/211
            PD   27-AUG-2002
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            WILLIAM I WOOD,AUSTIN L GURNEY,AUDREY GODDARD,DIANE PENNICA, PI
            JIAN ZHENG,
            PI   JEAN YUAN
            PC   C12N15/09,C07K14/47,C07K16/18,C07K19/00,C12N1/19,C12N1/21, PC
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            (C12N5/00,C12R1:91)
            CC   Secreted and transmembrane polypeptides and nucleic CC   acids
            encoding the same
            FH   Key             Location/Qualifiers
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            FT                   /organism='Homo sapiens (human)'.
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                     /mol_type="genomic DNA"
                     /db_xref="taxon:9606"
ORIGIN

  Query Match           99.1%;  Score 1090.6;  DB 6;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 1.5e-264;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
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Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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RESULT 4
BD173076
LOCUS       BD173076                1100 bp    DNA     linear   PAT 18-FEB-2003
DEFINITION  Secreted and transmembrane polypeptides and nucleic acids encoding
            the same.
ACCESSION   BD173076
VERSION     BD173076.1  GI:28414385
KEYWORDS    JP 2002238587-A/211.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 1100)
  AUTHORS   Wood,W.I., Gurney,A.L., Goddard,A., Pennica,D., Zheng,J. and
            Yuan,J.
  TITLE     Secreted and transmembrane polypeptides and nucleic acids encoding
            the same
  JOURNAL   Patent: JP 2002238587-A 211 27-AUG-2002;
            GENENTECH INC
COMMENT     OS   Homo sapiens (human)
            PN   JP 2002238587-A/211
            PD   27-AUG-2002
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            WILLIAM I WOOD,AUSTIN L GURNEY,AUDREY GODDARD,DIANE PENNICA, PI
            JIAN ZHENG,
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             (C12P21/02,C12R1:645),C12N15/00,C12N5/00,C12N15/00 CC   Secreted
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ORIGIN

  Query Match           99.1%;  Score 1090.6;  DB 6;  Length 1100;
  Best Local Similarity  99.6%;  Pred. No. 1.5e-264;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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Qy       1081 AAAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAAA 1100
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RESULT 5
BD173395
LOCUS BD173395 1100 bp DNA linear PAT 18-FEB-2003

DEFINITION Secreted and transmembrane polypeptides and nucleic acids encoding the same.
ACCESSION BD173395
VERSION BD173395.1 GI:28414706
KEYWORDS JP 2002238588-A/211.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 1100)
AUTHORS Wood,W.I., Gurney,A.L., Goddard,A., Pennica,D., Zheng,J. and Yuan,J.
TITLE Secreted and transmembrane polypeptides and nucleic acids encoding the same
JOURNAL Patent: JP 2002238588-A 211 27-AUG-2002;
GENENTECH INC
COMMENT OS Homo sapiens (human)
PN JP 2002238588-A/211
PD 27-AUG-2002
PF 18-DEC-2001 JP 2001385315
PR 17-SEP-1997 US 60/059115,17-SEP-1997 US 60/059184 PR
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WILLIAM I WOOD,AUSTIN L GURNEY,AUDREY GODDARD,DIANE PENNICA, PI JIAN ZHENG,
PI JEAN YUAN
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PC (C12N5/10,C12R1:91),C12N15/00,C12N5/00,(C12N5/00,C12R1:91) CC
Secreted and transmembrane polypeptides and nucleic CC acids encoding the same
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Query Match 99.1%; Score 1090.6; DB 6; Length 1100;
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Matches 1093; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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Qy         661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Qy        1081 AAAAAAAAAAAAAAAA 1097
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Db        1084 AAAAAAAAAAAAAAAA 1100


RESULT 6
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BD175429
LOCUS       BD175429               1100 bp    DNA     linear   PAT 18-MAR-2003
DEFINITION  Secretory and transmembrane polypeptide and nucleic acid encoding
            the same.
ACCESSION   BD175429
VERSION     BD175429.1  GI:29121127
KEYWORDS    JP 2002253280-A/211.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 1100)
  AUTHORS   Wood,W.I., Gurney,A.L., Goddard,A., Pennica,D., Zheng,J. and
            Yuan,J.
  TITLE     Secretory and transmembrane polypeptide and nucleic acid encoding
            the same
  JOURNAL   Patent: JP 2002253280-A 211 10-SEP-2002;
            GENENTECH INC
COMMENT     OS   Homo sapiens (human)
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            PD   10-SEP-2002
            PF   18-DEC-2001 JP 2001385319
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            WILLIAM I WOOD,AUSTIN L GURNEY,AUDREY GODDARD,DIANE PENNICA, PI
            JIAN ZHENG,
            PI   JEAN YUAN
            PC   C12N15/09,A61K45/00,A61P1/00,A61P13/12,A61P17/00,A61P17/06, PC
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ORIGIN

  Query Match           99.1%;  Score 1090.6;  DB 6;  Length 1100;
  Best Local Similarity 99.6%;  Pred. No. 1.5e-264;
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Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

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RESULT 7
AR410808
LOCUS       AR410808               1100 bp    DNA     linear   PAT 18-DEC-2003
DEFINITION  Sequence 256 from patent US 6635468.
ACCESSION   AR410808
VERSION     AR410808.1  GI:40162308
KEYWORDS    .
SOURCE      Unknown.
  ORGANISM  Unknown.
            Unclassified.
REFERENCE   1  (bases 1 to 1100)
  AUTHORS   Ashkenazi,A., Botstein,D., Desnoyers,L., Eaton,D.L., Ferrara,N.,
            Filvaroff,E., Fong,S., Gao,W.-Q., Gerber,H., Gerritsen,M.E.,
            Goddard,A., Godowski,P.J., Grimaldi,J.C., Gurney,A.L., Hillan,K.J.,
            Kljavin,I.J., Mather,J.P., Pan,J., Paoni,N.F., Roy,M.A.,
            Stewart,T.A., Tumas,D., Williams,P.M. and Wood,W.I.
  TITLE     Secreted and transmembrane polypeptides and nucleic acids encoding
            the same
  JOURNAL   Patent: US 6635468-A 256 21-OCT-2003;
FEATURES             Location/Qualifiers
     source          1. .1100
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ORIGIN

  Query Match             99.1%;  Score 1090.6;  DB 6;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 1.5e-264;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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RESULT 8
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LOCUS       AR439172                1100 bp    DNA     linear   PAT 20-FEB-2004
DEFINITION  Sequence 256 from patent US 6664376.
ACCESSION   AR439172
VERSION     AR439172.1  GI:42665021
KEYWORDS    .
SOURCE      Unknown.
  ORGANISM  Unknown.
            Unclassified.
REFERENCE   1  (bases 1 to 1100)
  AUTHORS   Ashkenazi,A., Botstein,D., Desnoyers,L., Eaton,D.L., Ferrara,N.,
            Filvaroff,E., Fong,S., Gao,W.-Q., Gerber,H., Gerritsen,M.E.,
            Goddard,A., Godowski,P.J., Grimaldi,J.C., Gurney,A.L., Hillan,K.J.,
            Kljavin,I.J., Mather,J.P., Pan,J., Paoni,N.F., Roy,M.A.,
            Stewart,T.A., Tumas,D., Williams,P.M. and Wood,W.I.
  TITLE     Secreted and transmembrane polypeptides and nucleic acids encoding
            the same
  JOURNAL   Patent: US 6664376-A 256 16-DEC-2003;
FEATURES             Location/Qualifiers
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ORIGIN

  Query Match             99.1%;  Score 1090.6;  DB 6;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 1.5e-264;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
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Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 9
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AR473192
LOCUS       AR473192               1100 bp    DNA     linear   PAT 20-FEB-2004
DEFINITION  Sequence 256 from patent US 6686451.
ACCESSION   AR473192
VERSION     AR473192.1  GI:42708567
KEYWORDS    .
SOURCE      Unknown.
  ORGANISM  Unknown.
            Unclassified.
REFERENCE   1  (bases 1 to 1100)
  AUTHORS   Desnoyers,L., Goddard,A., Godowski,P.J., Gurney,A.L., Mather,J.P.,
            Williams,P.M. and Wood,W.I.
  TITLE     Secreted and transmembrane polypeptides and nucleic acids encoding
            the same
  JOURNAL   Patent: US 6686451-A 256 03-FEB-2004;
FEATURES             Location/Qualifiers
     source          1. .1100
                     /organism="unknown"
                     /mol_type="genomic DNA"
ORIGIN

  Query Match           99.1%;  Score 1090.6;  DB 6;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 1.5e-264;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
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Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              ||||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 10
AR527178
LOCUS       AR527178               1100 bp    DNA     linear   PAT 08-OCT-2004
DEFINITION  Sequence 256 from patent US 6723535.
ACCESSION   AR527178
VERSION     AR527178.1  GI:53914095
KEYWORDS    .
SOURCE      Unknown.
  ORGANISM  Unknown.
            Unclassified.
REFERENCE   1  (bases 1 to 1100)
  AUTHORS   Ashkenazi,A., Botstein,D., Desnoyers,L., Eaton,D.L., Ferrara,N.,
            Filvaroff,E., Fong,S., Gao,W.-Q., Gerber,H., Gerritsen,M.E.,
            Goddard,A., Godowski,P.J., Grimaldi,J.C., Gurney,A.L., Hillan,K.J.,
            Kljavin,I.J., Mather,J.P., Pan,J., Paoni,N.F., Roy,M.A.,
            Stewart,T.A., Tumas,D., Williams,P.M. and Wood,W.I.
  TITLE     Secreted and transmembrane polypeptides and nucleic acids encoding
            the same
  JOURNAL   Patent: US 6723535-A 256 20-APR-2004;
FEATURES             Location/Qualifiers
     source          1. .1100
                     /organism="unknown"
                     /mol_type="genomic DNA"
ORIGIN

  Query Match           99.1%;  Score 1090.6;  DB 6;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 1.5e-264;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
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Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
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Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
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Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
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Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
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Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              |||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAAA 1100
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RESULT 11
AR566211
LOCUS       AR566211               1100 bp    DNA     linear   PAT 08-OCT-2004
DEFINITION  Sequence 256 from patent US 6767995.
ACCESSION   AR566211
VERSION     AR566211.1  GI:53983121
KEYWORDS    .
SOURCE      Unknown.
  ORGANISM  Unknown.
            Unclassified.
REFERENCE   1  (bases 1 to 1100)
  AUTHORS   Desnoyers,L., Goddard,A., Godowski,P.J., Gurney,A.L. and Wood,W.I.
  TITLE     Secreted and transmembrane polypeptides and nucleic acids encoding
            the same
  JOURNAL   Patent: US 6767995-A 256 27-JUL-2004;
FEATURES             Location/Qualifiers
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Query Match 99.1%; Score 1090.6; DB 6; Length 1100;
Best Local Similarity 99.6%; Pred. No. 1.5e-264;
Matches 1093; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
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Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
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Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

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Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

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Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
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Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              ||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903
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Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
              |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 12
AX697665
LOCUS       AX697665                1100 bp    DNA     linear   PAT 02-APR-2003
DEFINITION  Sequence 256 from Patent WO0104311.
ACCESSION   AX697665
VERSION     AX697665.1  GI:29498754
KEYWORDS    .
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1
  AUTHORS   Ashkenazi,A.J., Botstein,D., Desnoyers,L., Eaton,D.L., Ferrara,N.,
            Filvaroff,E., Fong,S., Gao,W.Q., Gerber,H., Gerritsen,M.E.,
            Goddard,A., Godowski,P.J., Grimaldi,C.J., Gurney,A.L., Hillan,K.J.,
            Kljavin,I.J., Mather,J.P., Pan,J., Paoni,N.F., Roy,M.A.,
            Stewart,T.A., Tumas,D., Williams,P.M. and Wood,W.I.
  TITLE     Secreted and transmembrane polypeptides and nucleic acids encoding
            the same
  JOURNAL   Patent: WO 0104311-A 256 18-JAN-2001;
            Genentech Inc. (US)
FEATURES             Location/Qualifiers
     source          1. .1100
                     /organism="Homo sapiens"
                     /mol_type="unassigned DNA"
                     /db_xref="taxon:9606"
ORIGIN

  Query Match             99.1%;  Score 1090.6;  DB 6;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 1.5e-264;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
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Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
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Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
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RESULT 13
BD075578
LOCUS       BD075578                1100 bp    DNA     linear   PAT 27-AUG-2002
DEFINITION  Secretory and transmembrane polypeptide and nucleic acid encoding
            the same.
ACCESSION   BD075578
VERSION     BD075578.1  GI:22621181
KEYWORDS    JP 2001516580-A/211.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 1100)
  AUTHORS   Wood,W.I., Gurney,A.L., Goddard,A., Penica,D., Chen,J. and Yuan,J.
  TITLE     Secretory and transmembrane polypeptide and nucleic acid encoding
            the same
  JOURNAL   Patent: JP 2001516580-A 211 02-OCT-2001;
            GENENTECH INC
COMMENT     OS   Homo sapiens (human)
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          PI   WILLIAM I WOOD,AUSTIN L GURNEY,AUDLEY GODDARD,DIANE PENICA, PI
            JEAN CHEN,
          PI   JEAN YUAN
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          PC   C12N15/00,C12N5/00
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FEATURES             Location/Qualifiers
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ORIGIN

  Query Match           99.1%;  Score 1090.6;  DB 6;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 1.5e-264;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
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Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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RESULT 14
AY359034
LOCUS       AY359034               1100 bp    mRNA    linear   PRI 03-OCT-2003
DEFINITION  Homo sapiens clone DNA42551 PRSS21 (UNQ266) mRNA, complete cds.
ACCESSION   AY359034
VERSION     AY359034.1  GI:37183185
KEYWORDS    FLI_CDNA.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 1100)
  AUTHORS   Clark,H.F., Gurney,A.L., Abaya,E., Baker,K., Baldwin,D., Brush,J.,
            Chen,J., Chow,B., Chui,C., Crowley,C., Currell,B., Deuel,B.,
            Dowd,P., Eaton,D., Foster,J., Grimaldi,C., Gu,Q., Hass,P.E.,
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            Lee,J., Lewis,L., Liao,D., Mark,M., Robbie,E., Sanchez,C.,
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            Schoenfeld,J., Seshagiri,S., Simmons,L., Singh,J., Smith,V.,
            Stinson,J., Vagts,A., Vandlen,R., Watanabe,C., Wieand,D., Woods,K.,
            Xie,M.H., Yansura,D., Yi,S., Yu,G., Yuan,J., Zhang,M., Zhang,Z.,
            Goddard,A., Wood,W.I. and Godowski,P.
  TITLE     The Secreted Protein Discovery Initiative (SPDI), a Large-Scale
            Effort to Identify Novel Human Secreted and Transmembrane Proteins:
            A Bioinformatics Assessment
  JOURNAL   Genome Res. 13 (10), 2265-2270 (2003)
   PUBMED   12975309
REFERENCE   2  (bases 1 to 1100)
  AUTHORS   Clark,H.F.
  TITLE     Direct Submission
  JOURNAL   Submitted (01-AUG-2003) Department of Bioinformatics, Genentech,
            Inc., 1 DNA Way, South San Francisco, CA 94080, USA
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ORIGIN

  Query Match           99.1%;  Score 1090.6;  DB 9;  Length 1100;
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  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
              |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 15
AR253955
LOCUS       AR253955               1094 bp    DNA     linear   PAT 20-DEC-2002
DEFINITION  Sequence 3 from patent US 6479274.
ACCESSION   AR253955
VERSION     AR253955.1  GI:27302450
KEYWORDS    .
SOURCE      Unknown.
  ORGANISM  Unknown.
            Unclassified.
REFERENCE   1  (bases 1 to 1094)
  AUTHORS   Antalis,T.M. and Hooper,J.D.
  TITLE     DNA molecules encoding human HELA2 or testisin serine proteinases
  JOURNAL   Patent: US 6479274-A 3 12-NOV-2002;
FEATURES             Location/Qualifiers
     source          1. .1094
                     /organism="unknown"
                     /mol_type="genomic DNA"
ORIGIN

  Query Match           98.0%;  Score 1078;  DB 6;  Length 1094;
  Best Local Similarity  99.5%;  Pred. No. 2.3e-261;
  Matches 1094;  Conservative    0;  Mismatches    0;  Indels    6;  Gaps    1;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
```

```
Qy        61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
Db        61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120

Qy       121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
Db       121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180

Qy       181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
Db       181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240

Qy       241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
Db       241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAAC------TGACCTTAGTGATCCCTCCGG 294

Qy       301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
Db       295 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 354

Qy       361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
Db       355 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 414

Qy       421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
Db       415 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 474

Qy       481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
Db       475 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 534

Qy       541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
Db       535 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 594

Qy       601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
Db       595 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 654

Qy       661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
Db       655 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 714

Qy       721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
Db       715 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 774

Qy       781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
Db       775 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 834

Qy       841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
Db       835 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 894

Qy       901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
Db       895 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 954

Qy       961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
Db       955 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1014

Qy      1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
Db      1015 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1074

Qy      1081 AAAAAAAAAAAAAAAAAAAA 1100
Db      1075 AAAAAAAAAAAAAAAAAAAA 1094
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Search completed: March 5, 2005, 04:37:28
Job time : 5122.55 secs

GenCore version 5.1.6

OM protein - protein search, using sw model

Run on: February 27, 2005, 20:11:50 ; Search time 25.5815 Seconds
(without alignments)
1181.014 Million cell updates/sec

Title: US-10-040-647-6
Perfect score: 1728
Sequence: 1 MGARGALLLALLLARAGLRK.....PSWPLLFPPLLWALPLLGPV 314

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : PIR_79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	579.5	33.5	343	1	A57014	prostasin (EC 3.4.
2	523.5	30.3	270	2	S56160	mast cell tryptase
3	513.5	29.7	638	1	KQMSPL	plasma kallikrein
4	513	29.7	275	2	C35863	tryptase (EC 3.4.2
5	511	29.6	276	2	A38654	mast cell proteina
6	510	29.5	275	2	A35863	tryptase (EC 3.4.2
7	510	29.5	638	1	KQRTPL	plasma kallikrein
8	509	29.5	273	2	A47246	tryptase (EC 3.4.2
9	509	29.5	275	2	B35863	tryptase (EC 3.4.2
10	509	29.5	275	2	A32410	tryptase (EC 3.4.2
11	507	29.3	638	1	KQHUP	plasma kallikrein
12	499	28.9	274	2	JC4171	tryptase (EC 3.4.2
13	498.5	28.8	274	2	A45754	tryptase (EC 3.4.2
14	478.5	27.7	625	1	KFHU1	coagulation factor
15	476.5	27.6	417	1	S00845	hepsin (EC 3.4.21.
16	474.5	27.5	366	2	JE0105	testicular serine
17	457.5	26.5	415	1	A34170	acrosin (EC 3.4.21
18	457.5	26.5	436	2	JX0172	acrosin (EC 3.4.21
19	454.5	26.3	810	1	PLHU	plasmin (EC 3.4.21
20	454	26.3	855	2	JC7731	membrane-bound arg
21	453	26.2	455	2	A61545	plasmin (EC 3.4.21
22	451.5	26.1	237	2	S68702	tryptase (EC 3.4.2
23	446	25.8	416	1	S33777	hepsin (EC 3.4.21.
24	443.5	25.7	4548	1	S00657	apoprotein(a) (EC
25	443	25.6	1113	2	JE0315	low-density lipopr
26	441	25.5	1035	1	A43090	enteropeptidase (E
27	439.5	25.4	421	2	S29599	acrosin (EC 3.4.21
28	439	25.4	367	2	JE0104	testicular serine
29	438	25.3	1019	1	A56318	enteropeptidase (E
30	436.5	25.3	460	2	B61545	plasmin (EC 3.4.21
31	435	25.2	786	1	A47547	serine proteinease
32	435	25.2	810	2	B30848	plasmin (EC 3.4.21
33	434.5	25.1	418	2	A37344	acrosin (EC 3.4.21
34	434.5	25.1	812	1	PLMS	plasmin (EC 3.4.21
35	433	25.1	790	1	PLPG	plasmin (EC 3.4.21
36	432	25.0	421	1	S11674	acrosin (EC 3.4.21
37	432	25.0	812	1	PLBO	plasmin (EC 3.4.21
38	431.5	25.0	431	2	S47538	acrosin (EC 3.4.21
39	428.5	24.8	437	2	S18407	acrosin (EC 3.4.21
40	427.5	24.7	266	2	S54146	trypsin (EC 3.4.21
41	427	24.7	810	2	I46260	plasmin (EC 3.4.21
42	426	24.7	1420	2	A32869	apolipoprotein(a)
43	421	24.4	1524	2	T30337	polyprotein - Afri
44	416	24.1	761	2	JC5759	brain-specific ser
45	413	23.9	269	2	B32410	mastocytoma protei

ALIGNMENTS

RESULT 1
A57014
prostasin (EC 3.4.21.-) precursor - human
C;Species: Homo sapiens (man)
C;Date: 24-May-1996 #sequence_revision 24-May-1996 #text_change 09-Jul-2004
C;Accession: A57014; A54866
R;Yu, J.X.; Chao, L.; Chao, J.
J. Biol. Chem. 270, 13483-13489, 1995
A;Title: Molecular cloning, tissue-specific expression, and cellular localization of hume
A;Reference number: A57014; MUID:95286644; PMID:7768952
A;Accession: A57014
A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-343 <RES>
A;Cross-references: UNIPROT:Q16651; GB:L41351; NID:g862304; PIDN:AAC41759.1; PID:g862305
A;Experimental source: prostate
A;Note: parts of this sequence were determined by protein sequencing
R;Yu, J.X.; Chao, L.; Chao, J.
J. Biol. Chem. 269, 18843-18848, 1994
A;Title: Prostasin is a novel human serine proteinase from seminal fluid. Purification, t
A;Reference number: A54866; MUID:94308140; PMID:8034638
A;Accession: A54866
A;Molecule type: protein
A;Residues: 45-64 <YUA>
C;Genetics:
A;Gene: GDB:PRSS8
A;Cross-references: GDB:676446; OMIM:600823
A;Map position: 16p11.2-16p11.2
C;Superfamily: trypsin; trypsin homology
C;Keywords: glycoprotein; hydrolase; serine proteinase; transmembrane protein
F;1-32/Domain: signal sequence #status predicted <SIG>
F;33-44,45-343/Product: prostasin #status predicted <MAT>
F;33-44/Domain: prostasin light chain #status predicted <CHL>
F;45-343/Domain: prostasin heavy chain #status predicted <CHH>
F;45-281/Domain: trypsin homology <TRY>
F;323-341/Domain: transmembrane #status predicted <TMM1>
F;37-154,70-86,168-244,201-223,234-262/Disulfide bonds: #status predicted
F;85,134,238/Active site: His, Asp, Ser #status predicted
F;159/Binding site: carbohydrate (Asn) (covalent) #status experimental

Query Match 33.5%; Score 579.5; DB 1; Length 343;
Best Local Similarity 41.0%; Pred. No. 2.8e-44;
Matches 141; Conservative 45; Mismatches 101; Indels 57; Gaps 12;

```
Qy          6 ALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRLWDSH 65
              |:|| | | |:|    |  ||     |||     :|| ||  |  |:|||| |:    |
Db         16 AILLYLGLLRSG-TGAEGAEA-----PCG-VAPQARITGGSSAVAGQWPWQVSITYEGVH 68

Qy         66 VCGVSLLSHRWALTAAHCFETYSDLSDPS-----GWMVQFG--QLTSMPSFWSLQAYYTR 118
              ||| ||:| :| |:|||||        ||      : |: |  || |    :|  |  :
Db         69 VCGGSLVSEQWVLSAAHCF--------PSEHHKEAYEVKLGAHQLDS----YSEDAKVST 116
```

```
Qy        119 YFVSNIYLSPRYL-GNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWG 177
Db        117 --LKDIIPHPSYLQEGSQGDIALLQLSRPITFSRYIRPICLPAANASFPNGLHCTVTGWG 174

Qy        178 YIKEDEALPSPHTLQEVQVAIINNSMCNHLF---LKYSFRKDIFGDMVCAGNAQGGKDAC 234
Db        175 HVAPSVSLLTPKPLQQLEVPLISRETCNCLYNIDAKPEEPHFVQEDMVCAGYVEGGKDAC 234

Qy        235 FGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQ--------KLMA 286
Db        235 QGDSGGPLSCPVEGLWYLTGIVSWGDACGARNRPGVYTLASSYASWIQSKVTELQPRVVP 294

Qy        287 QSGMSQPDPSW----------------PLLFFPLLWALPLLGP 313
Db        295 QTQESQPDSNLCGSHLAFSSAPAQGLLRPILFLPLGLALGLLSP 338
```

RESULT 2
S56160
mast cell tryptase precursor - Mongolian jird
C;Species: Meriones unguiculatus (Mongolian jird)
C;Date: 27-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 09-Jul-2004
C;Accession: S56160
R;Murakumo, Y.; Ide, H.; Itoh, H.; Tomita, M.; Kobayashi, T.; Maruyama, H.; Horii, Y.; N
Biochem. J. 309, 921-926, 1995
A;Title: Cloning of the cDNA encoding mast cell tryptase of Mongolian gerbil, Meriones u
A;Reference number: S56160; MUID:95366971; PMID:7639711
A;Accession: S56160
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-270 <MUR>
A;Cross-references: UNIPROT:P50342; EMBL:D31789; NID:g517122; PIDN:BAA06598.1; PID:g5171
C;Superfamily: trypsin; trypsin homology
F;26-262/Domain: trypsin homology <TRY>

  Query Match 30.3%; Score 523.5; DB 2; Length 270;
  Best Local Similarity 40.2%; Pred. No. 2.2e-39;
  Matches 117; Conservative 37; Mismatches 92; Indels 45; Gaps 9;

```
Qy          7 LLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR----LW 62
Db          5 LLLALPLFSLMHRSPLCQEWG--------------IVGGQEAPGNKWPWQVSLRANETYW 50

Qy         63 DSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF-- 120
Db         51 -RHFCGGSLIHPQWVLTAAHC--VGPTIADPNKVRVQ------------LRKQYLYYHDH 95

Qy        121 ---VSNIYLSPR-YLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGW 176
Db         96 LLAVSRIITHPTFYATQNGADIALLELKNPVNISSHVHPVSLPPASETFPSGTLCWVTGW 155

Qy        177 GYIKEDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKD----IFGDMVCAGNAQGGKD 232
Db        156 GNIDNDVSLPPPFPLKEVQVPVVENQLCDLKYHKGVYTGDNIHIVRDDMLCAGNE--GHD 213

Qy        233 ACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQK 283
Db        214 SCQGDSGGPLVCKVNGTWLQAGVVSWGEGCALPNRPGIYTRVTYYLDWIHR 264
```

RESULT 3
KQMSPL
plasma kallikrein (EC 3.4.21.34) precursor - mouse
C;Species: Mus musculus (house mouse)
C;Date: 30-Sep-1992 #sequence_revision 30-Sep-1992 #text_change 09-Jul-2004
C;Accession: A36557
R;Seidah, N.G.; Sawyer, N.; Hamelin, J.; Mion, P.; Beaubien, G.; Brachpapa, L.; Rochemon
DNA Cell Biol. 9, 737-748, 1990
A;Title: Mouse plasma kallikrein: cDNA structure, enzyme characterization, and comparisc
A;Reference number: A36557; MUID:91090844; PMID:2264928
A;Accession: A36557
A;Molecule type: mRNA
A;Residues: 1-638 <SEI>
A;Cross-references: UNIPROT:P26262; GB:M58588; NID:g200358; PIDN:AAA63393.1; PID:g200359
A;Note: part of this sequence, including the amino ends of both the heavy and light chair
C;Comment: This protein, synthesized in the liver, circulates as a noncovalent complex wi
C;Comment: The zymogen is activated by factor XIIa, which cleaves the molecule into a lig
 are linked by one or more disulfide bonds.
C;Superfamily: coagulation factor XI; trypsin homology
C;Keywords: blood coagulation; duplication; fibrinolysis; glycoprotein; hydrolase; inflan
F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-390/Product: plasma kallikrein heavy chain #status experimental <HCH>
F;20-109/Domain: apple repeat <AP1>
F;110-199/Domain: apple repeat <AP2>
F;200-289/Domain: apple repeat <AP3>
F;291-380/Domain: apple repeat <AP4>
F;391-638/Product: plasma kallikrein light chain #status experimental <LCH>
F;391-621/Domain: trypsin homology <TRY>
F;21-104,47-77,51-57,111-194,137-166,141-147,201-284,227-256,231-237,292-375,318-347,322-
F;127,215,308,396,494/Binding site: carbohydrate (Asn) (covalent) #status predicted
F;434,483,578/Active site: His, Asp, Ser #status predicted

  Query Match 29.7%; Score 513.5; DB 1; Length 638;
  Best Local Similarity 40.2%; Pred. No. 5e-38;
  Matches 101; Conservative 42; Mismatches 85; Indels 23; Gaps 8;

```
Qy         38 ITSRIVGGEDAELGRWPWQGSLR---LWDSHVCGVSLLSHRWALTAAHCFE--TYSDLSD 92
Db        387 INARIVGGTNASLGEWPWQVSLQVKLVSQTHLCGGSIIGRQWVLTAAHCFDGIPYPDVWR 446

Qy         93 PSGWMVQFGQLT-SMPSFWSLQAYYTRYFVSNIYLSPRY-LGNSPYDIALVKLSAPVTYT 150
Db        447 IYGGILSLSEITKETPS--------SR--IKELIIHQEYKVSEGNYDIALIKLQTPLNYT 496

Qy        151 KHIQPICLQASTFEFENRTDCWVTGWGYIKEDEALPSPHTLQEVQVAIINNSMCNHLFLK 210
Db        497 EFQKPICLPSKADTNTIYTNCWVTGWGYTKEQG--ETQNILQKATIPLVPNEECQKKYRD 554

Qy        211 YSFRKDIFGDMVCAGNAQGGKDACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGV 270
Db        555 YVINK----QMICAGYKEGGTDACKGDSGGPLVCKHSGRWQLVGITSWGEGCGRKDQPGV 610

Qy        271 YTNISHHFEWI 281
Db        611 YTKVSEYMDWI 621
```

RESULT 4
C35863
tryptase (EC 3.4.21.59) III precursor - human
C;Species: Homo sapiens (man)
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 07-Mar-2003
C;Accession: C35863; E35863; A38893
R;Vanderslice, P.; Ballinger, S.M.; Tam, E.K.; Goldstein, S.M.; Craik, C.S.; Caughey, G.H
Proc. Natl. Acad. Sci. U.S.A. 87, 3811-3815, 1990
A;Title: Human mast cell tryptase: multiple cDNAs and genes reveal a multigene serine pro
A;Reference number: A35863; MUID:90251647; PMID:2187193
A;Accession: C35863
A;Molecule type: mRNA
A;Residues: 9-275 <VAN>
A;Accession: E35863
A;Molecule type: DNA
A;Residues: 1-9 <VA2>
A;Cross-references: GB:M33494; NID:g3927804; PIDN:AAC83172.1; PID:g339977
A;Note: the first nine residues of this sequence are inferred from genomic DNA of tryptas
R;Vanderslice, P.
submitted to GenBank, April 1990
A;Reference number: A38893
A;Accession: A38893
A;Molecule type: mRNA
A;Residues: 9-131,'K',132-275 <VA3>
A;Cross-references: GB:M33493; NID:g339984; PIDN:AAA36780.1; PID:g339985
C;Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; serine proteinase; zymogen
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-30/Domain: activation peptide #status predicted <ACT>

F;31-275/Product: tryptase I #status predicted <MAT>
F;31-267/Domain: trypsin homology <TRY>
F;74,121,224/Active site: His, Asp, Ser #status predicted

Query Match 29.7%; Score 513; DB 2; Length 275;
Best Local Similarity 40.1%; Pred. No. 2e-38;
Matches 115; Conservative 43; Mismatches 93; Indels 36; Gaps 10;

```
Qy          7 LLLAL--LLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRLWD- 63
              ||||| | :||          ||| |   :||    ||||::|   :|||| |||: |
Db          5 LLLALPVLASRA--------YAAPAPGQALQRV---GIVGGQEAPRSKWPWQVSLRVRDR 53

Qy         64 --SHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF- 120
                | || ||:  :| ||||||     |: | :   ||  :           ||
Db         54 YWMHFCGGSLIHPQWVLTAAHC--VGPDVKDLAALRVQLRE---------QHLYYQDQLL 102

Qy        121 -VSNIYLSPR-YLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGY 178
               || | : |: |      ||||::|  ||  : |:  : |  ::  |     |||||||
Db        103 PVSRIIVHPQPYTAQIGADIALLELEEPVNVSSHVHTVTLPPASETFPPGMPCWVTGWGD 162

Qy        179 IKEDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKD----IFGDMVCAGNAQGGKDAC 234
              :  || || |  |::|:| |: | :|:  :   ::  |   :  ||:|||| :  :|:|
Db        163 VDNDERLPPPFPLKQVKVPIMENHICDAKYHLGAYTGDDVRIVRDDMLCAGNTR--RDSC 220

Qy        235 FGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWI 281
              ||||||| |  :| | | |||||| || :|||||:|| :::: :||
Db        221 QGDSGGPLVCKVNGTWLQAGVVSWGEGCAQPNRPGIYTRVTYYLDWI 267
```

RESULT 5
A38654
mast cell proteinase 6 (EC 3.4.21.-) precursor - mouse
C;Species: Mus musculus (house mouse)
C;Date: 21-Feb-1992 #sequence_revision 17-Feb-1994 #text_change 09-Jul-2004
C;Accession: A38654; B38654; D35646; I59478
R;Reynolds, D.S.; Gurley, D.S.; Austen, K.F.; Serafin, W.E.
J. Biol. Chem. 266, 3847-3853, 1991
A;Title: Cloning of the cDNA and gene of mouse mast cell protease-6. Transcription by pr
A;Reference number: A38654; MUID:91139682; PMID:1995638
A;Accession: A38654
A;Molecule type: DNA
A;Residues: 1-276 <REY>
A;Cross-references: UNIPROT:P21845; GB:M57625; NID:g200506; PIDN:AAA39987.1; PID:g200507
A;Note: the authors translated the codon CGC for residue 24 as Ala, GAG for residue 37 a
s Gly, GAG for residue 148 as Gly, GAG for residue 168 as Gly, and GAA for 185 as Gly
A;Accession: B38654
A;Molecule type: mRNA
A;Residues: 1-276 <RE2>
A;Cross-references: GB:M57626; NID:g200508; PIDN:AAA39988.1; PID:g200509
R;Reynolds, D.S.; Stevens, R.L.; Lane, W.S.; Carr, M.H.; Austen, K.F.; Serafin, W.E.
Proc. Natl. Acad. Sci. U.S.A. 87, 3230-3234, 1990
A;Title: Different mouse mast cell populations express various combinations of at least
A;Reference number: A35646; MUID:90222202; PMID:2326280
A;Accession: D35646
A;Molecule type: protein
A;Residues: 32-54 <RE3>
R;Huang, R.; Abrink, M.; Gobl, A.E.; Nilsson, G.; Aveskogh, M.; Larsson, L.G.; Nilsson,
Scand. J. Immunol. 38, 359-367, 1993
A;Title: Expression of a mast cell tryptase in the human monocytic cell lines U-937 and
A;Reference number: I59478; MUID:94023807; PMID:8210998
A;Accession: I59478
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-276 <RES>
A;Cross-references: GB:L31853; NID:g473480; PIDN:AAA39725.1; PID:g473481
C;Genetics:
A;Gene: MMCP-6
A;Introns: 24/1; 79/2; 168/1; 222/3
C;Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; serine proteinase; zymogen
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-31/Domain: activation peptide #status predicted <ACT>
F;32-276/Product: mast cell proteinase 6 #status experimental <MAT>

F;32-268/Domain: trypsin homology <TRY>
F;75,122,225/Active site: His, Asp, Ser #status predicted

Query Match 29.6%; Score 511; DB 2; Length 276;
Best Local Similarity 39.3%; Pred. No. 3e-38;
Matches 106; Conservative 42; Mismatches 88; Indels 34; Gaps 9;

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Qy         32 PCGRRVITSRIVGGEDAELGRWPWQGSLRL----WDSHVCGVSLLSHRWALTAAHCFETY 87
              |  :||    |||| :|   :|||| |||     |  | || ||:  :| ||||||   :
Db         25 PANQRV---GIVGGHEASESKWPWQVSLRFKLNYW-IHFCGGSLIHPQWVLTAAHCVGPH 80

Qy         88 SDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRY-----FVSNIYLSPR-YLGNSPYDIALV 141
                :  |  : ||             |:  |  |       :: | : |  |       |:||:
Db         81 --IKSPQLFRVQ------------LREQYLYYGDQLLSLNRIVVHPHYYTAEGGADVALL 126

Qy        142 KLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIKEDEALPSPHTLQEVQVAIINN 201
              :|  ||  : || || |  ::  |   | |||||| |  || || |: |::|:| |: |
Db        127 ELEVPVNVSTHIHPISLPPASETFPPGTSCWVTGWGDIDNDEPLPPPYPLKQVKVPIVEN 186

Qy        202 SMCNHLFLKYSFRKDIF----GDMVCAGNAQGGKDACFGDSGGPLACNKDGLWYQIGVVS 257
              |:|:  :    :  | |      |:|||| :  :|:| ||||||| |   | | | ||||
Db        187 SLCDRKYHTGLYTGDDFPIVHDGMLCAGNTR--RDSCQGDSGGPLVCKVKGTWLQAGVVS 244

Qy        258 WGVGCGRPNRPGVYTNISHHFEWIQKLMAQ 287
              || || :||:||:|| :::: :|| : : :
Db        245 WGEGCAQPNKPGIYTRVTYYLDWIHRYVPE 274
```

RESULT 6
A35863
tryptase (EC 3.4.21.59) I precursor - human
C;Species: Homo sapiens (man)
C;Date: 12-Oct-1990 #sequence_revision 12-Oct-1990 #text_change 09-Jul-2004
C;Accession: A35863; D35863; A60939; A39326
R;Vanderslice, P.; Ballinger, S.M.; Tam, E.K.; Goldstein, S.M.; Craik, C.S.; Caughey, G.H
Proc. Natl. Acad. Sci. U.S.A. 87, 3811-3815, 1990
A;Title: Human mast cell tryptase: multiple cDNAs and genes reveal a multigene serine pro
A;Reference number: A35863; MUID:90251647; PMID:2187193
A;Accession: A35863
A;Molecule type: DNA
A;Residues: 1-275 <VAN>
A;Cross-references: UNIPROT:Q15661; GB:M33494; NID:g3927804; PIDN:AAC83172.1; PID:g339977
A;Accession: D35863
A;Molecule type: mRNA
A;Residues: 1-275 <VA2>
A;Cross-references: GB:M33491
R;Butterfield, J.H.; Weiler, D.A.; Hunt, L.W.; Wynn, S.R.; Roche, P.C.
J. Leukoc. Biol. 47, 409-419, 1990
A;Title: Purification of tryptase from a human mast cell line.
A;Reference number: A60939; MUID:90244210; PMID:2110591
A;Accession: A60939
A;Molecule type: protein
A;Residues: 31-38,'P',40-41,'X',43,'T',45-48,'X',50 <BUT>
A;Experimental source: mast cell
A;Note: 44-Gly was also found
R;Cromlish, J.A.; Seidah, N.G.; Marcinkiewicz, M.; Hamelin, J.; Johnson, D.A.; Chretien,
J. Biol. Chem. 262, 1363-1373, 1987
A;Title: Human pituitary tryptase: molecular forms, NH-2-terminal sequence, immunocytoche
A;Reference number: A39326; MUID:87109258; PMID:3543004
A;Accession: A39326
A;Molecule type: protein
A;Residues: 31-38 <CRO>
A;Experimental source: pituitary
C;Genetics:
A;Introns: 21/1; 78/2; 177/1; 221/3
C;Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; serine proteinase; zymogen
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-30/Domain: activation peptide #status predicted <ACT>
F;31-275/Product: tryptase I #status experimental <MAT>
F;31-267/Domain: trypsin homology <TRY>
F;74,121,224/Active site: His, Asp, Ser #status predicted

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  Query Match             29.5%;  Score 510;  DB 2;  Length 275;
  Best Local Similarity   39.9%;  Pred. No. 3.7e-38;
  Matches  115;  Conservative   43;  Mismatches   92;  Indels   38;  Gaps   11;

Qy          7 LLLAL--LLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRL--- 61
              |||||  | :||          |||  |   :||    ||||::|   :|||| |||:
Db          5 LLLALPVLASRA---------YAAPAPGQALQRV---GIVGGQEAPRSKWPWQVSLRVHGP 53

Qy         62 -WDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
               |  | || ||:  :| ||||||     |: | :   ||  :             ||
Db         54 YW-MHFCGGSLIHPQWVLTAAHC--VGPDVKDLAALRVQLRE---------QHLYYQDQL 101

Qy        121 --VSNIYLSPR-YLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWG 177
                || | : |: |      ||||::|  ||  : |:  : |  ::  |     |||||||
Db        102 LPVSRIIVHPQFYTAQIGADIALLELEEPVNVSSHVHTVTLPPASETFPPGMPCWVTGWG 161

Qy        178 YIKEDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKD----IFGDMVCAGNAQGGKDA 233
               :  || || |  |::|:| |: | :|:  :   ::  |    :  ||:|||| :  :|:
Db        162 DVDNDERLPPPFPLKQVKVPIMENHICDAKYHLGAYTGDDVRIVRDDMLCAGNTR--RDS 219

Qy        234 CFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWI 281
               | ||||||| |  :| | | ||||||  || :|||||:|| :::: :||
Db        220 CQGDSGGPLVCKVNGTWLQAGVVSWGEGCAQPNRPGIYTRVTYYLDWI 267


RESULT 7
KQRTPL
plasma kallikrein (EC 3.4.21.34) precursor - rat
N;Alternate names: Fletcher factor; kininogenin; serum kallikrein
C;Species: Rattus norvegicus (Norway rat)
C;Date: 30-Sep-1992 #sequence_revision 30-Sep-1992 #text_change 09-Jul-2004
C;Accession: A39180; A33320; S06851; I53041; S06852
R;Beaubien, G.; Rosinski-Chupin, I.; Mattei, M.G.; Mbikay, M.; Chretien, M.; Seidah, N.G
Biochemistry 30, 1628-1635, 1991
A;Title: Gene structure and chromosomal localization of plasma kallikrein.
A;Reference number: A39180; MUID:91129236; PMID:1993180
A;Accession: A39180
A;Molecule type: DNA
A;Residues: 1-638 <BEA>
A;Cross-references: UNIPROT:P14272; GB:J05315
A;Note: the authors translated the codon GAG for residue 81 as Gln
R;Seidah, N.G.; Ladenheim, R.; Mbikay, M.; Hamelin, J.; Lutfalla, G.; Rougeon, F.; Lazur
DNA 8, 563-574, 1989
A;Title: The cDNA structure of rat plasma kallikrein.
A;Reference number: A33320; MUID:90091743; PMID:2598771
A;Accession: A33320
A;Status: not compared with conceptual translation
A;Molecule type: mRNA
A;Residues: 1-638 <SEI>
A;Cross-references: GB:M30282; NID:g205010; PIDN:AAA41463.1; PID:g205011
A;Note: part of this sequence, including the amino ends of both the heavy and light chai
R;Paquin, J.; Benjannet, S.; Sawyer, N.; Lazure, C.; Chretien, M.; Seidah, N.G.
Biochim. Biophys. Acta 999, 103-110, 1989
A;Title: Rat plasma kallikrein: purification, NH(2)-terminal sequencing and development
A;Reference number: S06851; MUID:90089457; PMID:2597701
A;Accession: S06851
A;Molecule type: protein
A;Residues: 20-45;391-413 <PAQ>
R;Seidah, N.G.; Ladenheim, R.; Mbikay, M.; Hamelin, J.; Lutfalla, G.; Rougeon, R.; Lazur
DNA Cell Biol. 8, 563-574, 1989
A;Title: The cDNA structure of rat plasma kallikrein.
A;Reference number: I53041
A;Accession: I53041
A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-638 <RES>
A;Cross-references: GB:M58590; NID:g206721; PIDN:AAA42069.1; PID:g206722
C;Comment: This protein, synthesized in the liver, circulates as a noncovalent complex w
C;Comment: The zymogen is activated by factor XIIa, which cleaves the molecule into a li
 are linked by one or more disulfide bonds.
C;Genetics:
A;Gene: PK
C;Superfamily: coagulation factor XI; trypsin homology
C;Keywords: blood coagulation; duplication; fibrinolysis; glycoprotein; hydrolase; inflam
F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-390/Product: plasma kallikrein heavy chain #status experimental <MAT1>
F;20-109/Domain: apple repeat <AP1>
F;110-199/Domain: apple repeat <AP2>
F;200-289/Domain: apple repeat <AP3>
F;291-380/Domain: apple repeat <AP4>
F;391-638/Product: plasma kallikrein light chain #status experimental <MAT2>
F;391-621/Domain: trypsin homology <TRY>
F;21-104,47-77,51-57,111-194,137-166,141-147,201-284,227-256,231-237,292-375,318-347,322-
F;127,215,308,453,459,494/Binding site: carbohydrate (Asn) (covalent) #status predicted
F;396/Binding site: carbohydrate (Asn) (covalent) #status experimental
F;434,483,578/Active site: His, Asp, Ser #status predicted

  Query Match             29.5%;  Score 510;  DB 1;  Length 638;
  Best Local Similarity   37.7%;  Pred. No. 1e-37;
  Matches  100;  Conservative   47;  Mismatches   96;  Indels   22;  Gaps    7;

Qy         30 SGPCGRRVITSRIVGGEDAELGRWPWQGSLR---LWDSHVCGVSLLSHRWALTAAHCFE- 85
              |  |  : |:||||| :: || |||| ||:   :  :|:|| |::  :| |||||||:
Db        380 SSDCTTK-INARIVGGTNSSLGEWPWQVSLQVKLVSQNHMCGGSIIGRQWILTAAHCFDG 438

Qy         86 -TYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYFVSNIYLSPRY-LGNSPYDIALVKL 143
                | |:    | ::   ::|:   | |          :  : :  :| :    |||||:||
Db        439 IPYPDVWRIYGGILNLSEITNKTPFSS---------IKELIIHQKYKMSEGSYDIALIKL 489

Qy        144 SAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIKEDEALPSPHTLQEVQVAIINNSM 203
               |: ||:  :|||| :        |:|||||||| ||      : : ||:  : :: |
Db        490 QTPLNYTEFQKPICLPSKADTNTIYTNCWVTGWGYTKERG--ETQNILQKATIPLVPNEE 547

Qy        204 CNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGGPLACNKDGLWYQIGVVSWGVGCG 263
              |   :  |   |     |:|||  :|| ||| |||||||  |   | |  :|: ||| ||
Db        548 CQKKYRDYVITK----QMICAGYKEGGIDACKGDSGGPLVCKHSGRWQLVGITSWGEGCA 603

Qy        264 RPNRPGVYTNISHHFEWIQKLMAQS 288
               |  :|||||  :: : :|| : :  |
Db        604 RKEQPGVYTKVAEYIDWILEKIQSS 628


RESULT 8
A47246
tryptase (EC 3.4.21.59) 2 - mouse
C;Species: Mus musculus (house mouse)
C;Date: 21-Sep-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C;Accession: A47246
R;McNeil, H.P.; Reynolds, D.S.; Schiller, V.; Ghildyal, N.; Gurley, D.S.; Austen, K.F.; S
Proc. Natl. Acad. Sci. U.S.A. 89, 11174-11178, 1992
A;Title: Isolation, characterization, and transcription of the gene encoding mouse mast c
A;Reference number: A47246; MUID:93087489; PMID:1454796
A;Accession: A47246
A;Status: preliminary
A;Molecule type: nucleic acid
A;Residues: 1-273 <MCN>
A;Cross-references: UNIPROT:Q02844; GB:L00653; NID:g200518; PIDN:AAA39992.1; PID:g200519
A;Note: sequence extracted from NCBI backbone (NCBIN:119745, NCBIP:119746)
C;Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; serine proteinase
F;29-265/Domain: trypsin homology <TRY>

  Query Match             29.5%;  Score 509;  DB 2;  Length 273;
  Best Local Similarity   42.4%;  Pred. No. 4.5e-38;
  Matches  114;  Conservative   35;  Mismatches   86;  Indels   34;  Gaps    9;

Qy         26 AAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRLWDS---HVCGVSLLSHRWALTAAH 82
              |||  ||   |     ||||::|   :|||| |||  |:   | || ||:  :| |||||
Db         18 AAP--GPAMTR---EGIVGGQEAHGNKWPWQVSLRANDTYWMHFCGGSLIHPQWVLTAAH 72

Qy         83 CFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF----VSNIYLSPR-YLGNSPY 136
              |     |::||:   ||             |:  |  |       || |   |  |:
Db         73 C--VGPDVADPNKVRVQ-----------LRKQYLYYHDHLMTVSQIITHPDFYIVQDGA 118

Qy        137 DIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIKEDEALPSPHTLQEVQV 196
              ||||:||: ||  : :: |: |  ::  | : | |||||||  |      || |  |:||||
```

```
Db        119 DIALLKLTNPVNISDYVHPVPLPPASETFPSGTLCWVTGWGNIDNGVNLPPPFPLKEVQV 178

Qy        197 AIINNSMCNHLFLKYSFRKD----IFGDMVCAGNAQGGKDACFGDSGGPLACNKDGLWYQ 252

Db        179 PIIENHLCDLKYHKGLITGDNVHIVRDDMLCAGNE--GHDSCQGDSGGPLVCKVEDTWLQ 236

Qy        253 IGVVSWGVGCGRPNRPGVYTNISHHFEWI 281

Db        237 AGVVSWGEGCAQPNRPGIYTRVTYYLDWI 265


RESULT 9
B35863
tryptase (EC 3.4.21.59) II precursor - human
N;Alternate names: tryptase beta
C;Species: Homo sapiens (man)
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 09-Jul-2004
C;Accession: B35863; A37193; I59473
R;Vanderslice, P.; Ballinger, S.M.; Tam, E.K.; Goldstein, S.M.; Craik, C.S.; Caughey, G.
Proc. Natl. Acad. Sci. U.S.A. 87, 3811-3815, 1990
A;Title: Human mast cell tryptase: multiple cDNAs and genes reveal a multigene serine pr
A;Reference number: A35863; MUID:90251647; PMID:2187193
A;Accession: B35863
A;Molecule type: mRNA; DNA
A;Residues: 1-275 <VAN>
A;Cross-references: UNIPROT:P20231; GB:M33492; NID:g339982; PIDN:AAA36779.1; PID:g339983
A;Note: residues 2-275 are derived from mRNA; residue one was inferred from the genomic
R;Miller, J.S.; Moxley, G.; Schwartz, L.B.
J. Clin. Invest. 86, 864-870, 1990
A;Title: Cloning and characterization of a second complementary DNA for human tryptase.
A;Reference number: A37193; MUID:90369005; PMID:2203827
A;Accession: A37193
A;Molecule type: mRNA
A;Residues: 1-275 <MIL>
A;Cross-references: GB:M37488; NID:g179583; PIDN:AAA51843.1; PID:g179584
R;Blom, T.; Hellman, L.
Scand. J. Immunol. 37, 203-208, 1993
A;Title: Characterization of a tryptase mRNA expressed in the human basophil cell line K
A;Reference number: I59473; MUID:93166209; PMID:8434231
A;Accession: I59473
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-275 <RES>
A;Cross-references: GB:S55551; NID:g265666; PIDN:AAD13876.1; PID:g4261576
A;Experimental source: basophil cell line KU812
C;Genetics:
A;Gene: GDB:TPS1
A;Cross-references: GDB:125890; OMIM:191080
A;Map position: 16pter-16qter
C;Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; serine proteinase; zymogen
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-30/Domain: activation peptide #status predicted <ACT>
F;31-275/Product: tryptase I #status predicted <MAT>
F;31-267/Domain: trypsin homology <TRY>
F;74,121,224/Active site: His, Asp, Ser #status predicted

  Query Match             29.5%;  Score 509;  DB 2;  Length 275;
  Best Local Similarity   39.9%;  Pred. No. 4.6e-38;
  Matches  115;  Conservative   43;  Mismatches   92;  Indels   38;  Gaps   11;

Qy          7 LLLAL--LLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRL--- 61

Db          5 LLLALPVLASRA--------YAAPAPGQALQRV---GIVGGQEAPRSKWPWQVSLRVHGP 53

Qy         62 -WDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Db         54 YW-MHFCGGSLIHPQWVLTAAHC--VGPDVKDLAALRVQLRE---------QHLYYQDQL 101

Qy        121 --VSNIYLSPR-YLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWG 177

Db        102 LPVSRIIVHPQFYTAQIGADIALLELEEPVKVSSHVHTVTLPPASETFPPGMPCWVTGWG 161

Qy        178 YIKEDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKD----IFGDMVCAGNAQGGKDA 233

Db        162 DVDNDERLPPPFPLKQVKVPIMENHICDAKYHLGAYTGDDVRIVRDDMLCAGNTR--RDS 219

Qy        234 CFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWI 281

Db        220 CQGDSGGPLVCKVNGTWLQAGVVSWGEGCAQPNRPGIYTRVTYYLDWI 267


RESULT 10
A32410
tryptase (EC 3.4.21.59) precursor - dog
C;Species: Canis lupus familiaris (dog)
C;Date: 12-Oct-1989 #sequence_revision 12-Oct-1989 #text_change 09-Jul-2004
C;Accession: A32410
R;Vanderslice, P.; Craik, C.S.; Nadel, J.A.; Caughey, G.H.
Biochemistry 28, 4148-4155, 1989
A;Title: Molecular cloning of dog mast cell tryptase and a related protease: structural e
A;Reference number: A32410; MUID:89352460; PMID:2504277
A;Accession: A32410
A;Molecule type: mRNA
A;Residues: 1-275 <VAN>
A;Cross-references: UNIPROT:P15944; GB:M24664; NID:g163982; PIDN:AAA30854.1; PID:g163983;
C;Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; serine proteinase; zymogen
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-30/Domain: activation peptide #status predicted <ACT>
F;31-275/Product: tryptase #status predicted <MAT>
F;31-267/Domain: trypsin homology <TRY>
F;74,121,224/Active site: His, Asp, Ser #status predicted

  Query Match             29.5%;  Score 509;  DB 2;  Length 275;
  Best Local Similarity   40.6%;  Pred. No. 4.6e-38;
  Matches  116;  Conservative   40;  Mismatches   98;  Indels   32;  Gaps   11;

Qy          9 LALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRL----WDS 64

Db          5 LVLALALLGSLVPVS----PAPGQALQRV---GIVGGREAPGSKWPWQVSLRLKGQYW-R 56

Qy         65 HVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF--VS 122

Db         57 HICGGSLIHPQWVLTAAHC--VGPNVVCPEEIRVQLRE---------QHLYYQDHLLPVN 105

Qy        123 NIYLSPR-YLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIKE 181

Db        106 RIVMHPNYYTPENGADIALLELEDPVNVSAHVQPVTLPPALQTFPTGTPCWVTGWGDVHS 165

Qy        182 DEALPSPHTLQEVQVAIINNSMCN---HLFLKYSFR-KDIFGDMVCAGNAQGGKDACFGD 237

Db        166 GTPLPPPFPLKQVKVPIVENSMCDVQYHLGLSTGDGVRIVREDMLCAGNSK--SDSCQGD 223

Qy        238 SGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQK 283

Db        224 SGGPLVCRVRGVWLQAGVVSWGEGCAQPNRPGIYTRVAYYLDWIHQ 269


RESULT 11
KQHUP
plasma kallikrein (EC 3.4.21.34) precursor - human
N;Alternate names: kininogenin; plasma prekallikrein
C;Species: Homo sapiens (man)
C;Date: 13-Aug-1986 #sequence_revision 13-Aug-1986 #text_change 09-Jul-2004
C;Accession: A00921; A37939
R;Chung, D.W.; Fujikawa, K.; McMullen, B.A.; Davie, E.W.
Biochemistry 25, 2410-2417, 1986
A;Title: Human plasma prekallikrein, a zymogen to a serine protease that contains four ta
A;Reference number: A00921; MUID:86243359; PMID:3521732
A;Accession: A00921
A;Molecule type: mRNA
A;Residues: 1-638 <CHU>
A;Cross-references: UNIPROT:P03952; GB:M13143; NID:g190262; PIDN:AAA60153.1; PID:g190263
R;McMullen, B.A.; Fujikawa, K.; Davie, E.W.
Biochemistry 30, 2050-2056, 1991
```

```
A;Title: Location of the disulfide bonds in human plasma prekallikrein: the presence of
A;Reference number: A37939; MUID:91152016; PMID:1998666
A;Accession: A37939
A;Molecule type: protein
A;Residues: 20-27;40-46,'X',48,'H';50,'X',52-70,'H';75-76,'X',78-80;103-113;131-140;141-
;260-283,'X',285;287-291,'X',293-295;314-317,'X',319-320;321-324;'X',329-333;334-339,'X'
525;538-551;562,'X',564-567;573,'X',575-576;578-583,'X',585;592-604 <MCM>
C;Comment: This protein, synthesized in the liver, circulates as a noncovalent complex w
C;Comment: The zymogen is activated by factor XIIa, which cleaves the molecule into a li
 are linked by one or more disulfide bonds.
C;Comment: The enzyme cleaves Lys-Arg and Arg-Ser bonds. It activates, in a reciprocal r
inogen and may also play a role in the renin-angiotensin system by converting prorenin i
C;Genetics:
A;Gene: GDB:KLK3
A;Cross-references: GDB:127575; OMIM:229000
A;Map position: 4q35-4q35
C;Superfamily: coagulation factor XI; trypsin homology
C;Keywords: blood coagulation; duplication; fibrinolysis; glycoprotein; hydrolase; infla
F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-638/Product: plasma kallikrein #status predicted <MAT>
F;20-390/Domain: plasma kallikrein heavy chain #status predicted <HCH>
F;20-109/Domain: apple repeat <AP1>
F;110-199/Domain: apple repeat <AP2>
F;200-289/Domain: apple repeat <AP3>
F;291-380/Domain: apple repeat <AP4>
F;391-638/Domain: plasma kallikrein light chain #status predicted <LCH>
F;391-621/Domain: trypsin homology <TRY>
F;21-104,47-77,51-57,111-194,137-166,141-147,201-284,227-256,231-237,292-375,322-328,383
F;127,308,396,453,494/Binding site: carbohydrate (Asn) (covalent) #status experimental
F;318-347,340-345/Disulfide bonds: #status predicted
F;390-391/Cleavage site: Arg-Ile (coagulation factor XIIa) #status predicted
F;434,483,578/Active site: His, Asp, Ser #status predicted

  Query Match             29.3%;  Score 507;  DB 1;  Length 638;
  Best Local Similarity   39.6%;  Pred. No. 1.9e-37;
  Matches  105;  Conservative   43;  Mismatches   87;  Indels   30;  Gaps    9;

Qy         39 TSRIVGGEDAELGRWPWQGSLRL---WDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSG 95
              ::|||||  ::  |  |||| ||::      |:|| ||: |:| |||||||:    |
Db        388 STRIVGGTNSSWGEWPWQVSLQVKLTAQRHLCGGSLIGHQWVLTAAHCFD---GLPLQDV 444

Qy         96 WMVQFG-----QLTSMPSFWSLQ--AYYTRYFVSNIYLSPRYLGNSPYDIALVKLSAPVT 148
              | :  |      :|    |  ::    :  | ||          ||  :||||:|| ||:
Db        445 WRIYSGILNLSDITKDTPFSQIKEIIIHQNYKVSE--------GN--HDIALIKLQAPLN 494

Qy        149 YTKHIQPICLQASTFEFENRTDCWVTGWGYIKEDEALPSPHTLQEVQVAIINNSMCNHLF 208
              ||:  :|||| :       |:|||||||: ||   :   : ||:| : :: |  |   :
Db        495 YTEFQKPICLPSKGDTSTIYTNCWVTGWGFSKEKGEI--QNILQKVNIPLVTNEECQKRY 552

Qy        209 LKYSFRKDIFGDMVCAGNAQGGKDACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRP 268
                |     |   |||||  :|||||| ||||||| |  :|:|  :|: ||| || |  :|
Db        553 QDYK----ITQRMVCAGYKEGGKDACKGDSGGPLVCKHNGMWRLVGITSWGEGCARREQP 608

Qy        269 GVYTNISHHFEWI-QKLMAQSGMSQ 292
              |||| :: : :|| :|  :  | :|
Db        609 GVYTKVAEYMDWILEKTQSSDGKAQ 633


RESULT 12
JC4171
tryptase (EC 3.4.21.59) precursor - rat
N;Alternate names: mast cell tryptase
C;Species: Rattus norvegicus (Norway rat)
C;Date: 27-Aug-1995 #sequence_revision 27-Oct-1995 #text_change 09-Jul-2004
C;Accession: JC4171
R;Ide, H.; Itoh, H.; Tomita, M.; Murakumo, Y.; Kobayashi, T.; Maruyama, H.; Osada, Y.; N
J. Biochem. 118, 210-215, 1995
A;Title: cDNA sequencing and expression of rat mast cell tryptase.
A;Reference number: JC4171; MUID:96015171; PMID:8537314
A;Accession: JC4171
A;Molecule type: mRNA
A;Residues: 1-274 <IDE>
A;Cross-references: UNIPROT:P50343; DDBJ:D38455; NID:g556555; PIDN:BAA07486.1; PID:g5565
C;Comment: This enzyme is basically specific for a connective tissue mast cell, it is up-
einase inhibitors.
C;Keywords: glycoprotein; hydrolase; mast cell; serine proteinase; zymogen
F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-29/Domain: activation peptide #status predicted <ACT>
F;30-274/Product: mast cell tryptase #status predicted <MAT>
F;30-266/Domain: trypsin homology <TRY>
F;73,120,223/Active site: His, Asp, Ser #status predicted
F;131/Binding site: carbohydrate (Asn) (covalent) #status predicted

  Query Match             28.9%;  Score 499;  DB 2;  Length 274;
  Best Local Similarity   39.3%;  Pred. No. 3.6e-37;
  Matches  116;  Conservative   39;  Mismatches  102;  Indels   38;  Gaps   11;

Qy          7 LLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR----LW 62
              | | |||| : |       |||   | :||    |||| :|   :|||| |||     |
Db          2 LKLLLLLALSPL--ASLVHAAPC--PVKQRV---GIVGGREASESKWPWQVSLRFKFSFW 54

Qy         63 DSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRY--- 119
                | || ||:  :| ||||||   :  :  |  : ||            |:  |  |
Db         55 -MHFCGGSLIHPQWVLTAAHCVGLH--IKSPELFRVQ-----------LREQYLYYADQ 99

Qy        120 --FVSNIYLSPRYLG-NSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGW 176
                 |:   : | |        ||||::|  ||  : || || |  ::  | : | ||||||
Db        100 LLTVNRTVVHPHYYTVEDGADIALLELEIPVNVSTHIHPISLPPASETFPSGTSCWVTGW 159

Qy        177 GYIKEDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKD----IFGDMVCAGNAQGGKD 232
              | |  || |  |: |::|:|  |: ||:|:  :    :  |    :   |:|||| :   |
Db        160 GDIDSDEPLLPPYPLKQVKVPIVENSLCDRKYHTGLYTGDDVPIVQDGMLCAGNTR--SD 217

Qy        233 ACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQ 287
              :| ||||||| |   | | | |||||| ||   ||||:|| :::: :|| : : |
Db        218 SCQGDSGGPLVCKVKGTWLQAGVVSWGEGCAEANRPGIYTRVTYYLDWIHRYVPQ 272


RESULT 13
A45754
tryptase (EC 3.4.21.59) alpha precursor - human
C;Species: Homo sapiens (man)
C;Date: 03-Jun-1993 #sequence_revision 03-Jun-1993 #text_change 08-Sep-1997
C;Accession: A45754; B37193
R;Miller, J.S.; Westin, E.H.; Schwartz, L.B.
J. Clin. Invest. 84, 1188-1195, 1989
A;Title: Cloning and characterization of complementary DNA for human tryptase.
A;Reference number: A45754; MUID:90009311; PMID:2677049
A;Accession: A45754
A;Molecule type: mRNA
A;Residues: 1-274 <MIL>
A;Cross-references: GB:M30038
R;Miller, J.S.; Moxley, G.; Schwartz, L.B.
J. Clin. Invest. 86, 864-870, 1990
A;Title: Cloning and characterization of a second complementary DNA for human tryptase.
A;Reference number: A37193; MUID:90369005; PMID:2203827
A;Accession: B37193
A;Molecule type: mRNA
A;Residues: 1-274 <MI2>
A;Cross-references: GB:M30038
A;Note: the sequence from Fig. 4 is inconsistent with that from Fig. 2 in having 205-Pro
C;Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; serine proteinase; zymogen
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-30/Domain: activation peptide #status predicted <ACT>
F;31-274/Product: tryptase I #status predicted <MAT>
F;31-266/Domain: trypsin homology <TRY>
F;74,120,223/Active site: His, Asp, Ser #status predicted

  Query Match             28.8%;  Score 498.5;  DB 2;  Length 274;
  Best Local Similarity   38.8%;  Pred. No. 4e-37;
  Matches  111;  Conservative   44;  Mismatches   96;  Indels   35;  Gaps    9;

Qy          7 LLLAL-LLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRLWD-- 63
              ||||| :||      |   :|   :|          ||||::|   :|||| |||: |
```

```
Db          5 LLLALPVLASRAYAAPAPVQALQQAG----------IVGGQEAPRSKWPWQVSLRVRDRY 54

Qy         64 -SHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF-- 120
                 | || ||:  :| ||||||     |: | :   | |             ||
Db         55 WMHFCGGSLIHPQWVLTAAHCLG--PDVKDLATLRVNSG----------THLYYQDQLLP 102

Qy        121 VSNIYLSPR-YLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYI 179
                || | : |: |:  :  ||||::|  ||  :  :  : |  ::  |     ||||||  :
Db        103 VSRIMVHPQFYIIQTGADIALLELEEPVNISSRVHTVMLPPASETFPPGMPCWVTGWGDV 162

Qy        180 KEDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKD----IFGDMVCAGNAQGGKDACF 235
                 || || |  |::|:|  |: | :|:  :   ::  |    |  ||:||||:|  :|:|
Db        163 DNDEPLPPPFPLKQVKVPIMENHICDAKYHLGAYTGDDVRIIRDDMLCAGNSQ--RDSCK 220

Qy        236 GDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWI 281
              |||||||  |  :| | | |||||  || :|||||:|| :::: :||
Db        221 GDSGGPLVCKVNGTWLQAGVVSWDEGCAQPNRPGIYTRVTYYLDWI 266


RESULT 14
KFHU1
coagulation factor XIa (EC 3.4.21.27) precursor [validated] - human
N;Alternate names: antihemophilic factor C; plasma thromboplastin antecedent
C;Species: Homo sapiens (man)
C;Date: 13-Aug-1986 #sequence_revision 26-May-1994 #text_change 09-Jul-2004
C;Accession: A27431; A00920; A37940
R;Asakai, R.; Davie, E.W.; Chung, D.W.
Biochemistry 26, 7221-7228, 1987
A;Title: Organization of the gene for human factor XI.
A;Reference number: A27431; MUID:88107663; PMID:2827746
A;Accession: A27431
A;Molecule type: DNA
A;Residues: 1-625 <ASA>
A;Cross-references: UNIPROT:P03951; GB:M18295
A;Note: the sequence shown follows the authors' translation
R;Fujikawa, K.; Chung, D.W.; Hendrickson, L.E.; Davie, E.W.
Biochemistry 25, 2417-2424, 1986
A;Title: Amino acid sequence of human factor XI, a blood coagulation factor with four ta
A;Reference number: A00920; MUID:86243360; PMID:3636155
A;Accession: A00920
A;Molecule type: mRNA
A;Residues: 1-625 <FUJ>
A;Cross-references: GB:M13142; NID:g182832; PIDN:AAA52487.1; PID:g182833
R;McMullen, B.A.; Fujikawa, K.; Davie, E.W.
Biochemistry 30, 2056-2060, 1991
A;Title: Location of the disulfide bonds in human coagulation factor XI: the presence of
A;Reference number: A37940; MUID:91152017; PMID:1998667
A;Accession: A37940
A;Molecule type: protein
A;Residues: 28-33;35-49,'X',51-55,'X',57-63;70-75,'X',77-79;107-109,'X',111-112;132-139;
;280-282,'X',284;285-297;313-316,'X',318-319;320-326;'X',328-330;'X',347-349;373,'X',375
C;Comment: The proenzyme consists of two identical chains linked by one or more disulfid
he active site, and a heavy chain, which associates with high molecular weight (HMW) kin
C;Genetics:
A;Gene: GDB:F11
A;Cross-references: GDB:119891; OMIM:264900
A;Map position: 4q35-4q35
A;Introns: 19/1; 73/2; 109/1; 162/2; 199/1; 252/2; 289/1; 343/2; 379/1; 435/2; 494/1; 52
C;Function:
A;Description: catalyzes the proteolytic activation of coagulation factor IX
A;Pathway: blood coagulation intrinsic pathway
C;Superfamily: coagulation factor XI; trypsin homology
C;Keywords: blood coagulation; duplication; glycoprotein; hemophilia C; homodimer; hydro
F;1-18/Domain: signal sequence #status predicted <SIG>
F;19-387/Product: coagulation factor XIa heavy chain #status experimental <HCH>
F;19-108/Domain: apple repeat <AP1>
F;109-198/Domain: apple repeat <AP2>
F;199-288/Domain: apple repeat <AP3>
F;290-379/Domain: apple repeat <AP4>
F;388-625/Product: coagulation factor XIa light chain #status experimental <LCH>
F;388-618/Domain: trypsin homology <TRY>
F;20-103,514-581,571-599/Disulfide bonds: #status predicted
F;29/Disulfide bonds: interchain #status experimental
F;46-76,50-56,110-193,136-165,140-146,200-283,226-255,230-236,291-374,317-346,321-327,38(
F;90,126,353,450/Binding site: carbohydrate (Asn) (covalent) #status predicted
F;339/Disulfide bonds: interchain #status predicted
F;387-388/Cleavage site: Arg-Ile (coagulation factor XIIa) #status experimental
F;431,480,575/Active site: His, Asp, Ser #status predicted
F;491/Binding site: carbohydrate (Asn) (covalent) #status experimental

  Query Match             27.7%;  Score 478.5;  DB 1;  Length 625;
  Best Local Similarity   40.6%;  Pred. No. 6.7e-35;
  Matches  102;  Conservative   40;  Mismatches   86;  Indels   23;  Gaps    8;

Qy         38 ITSRIVGGEDAELGRWPWQGSLRL---WDSHVCGVSLLSHRWALTAAHCF---ETYSDLS 91
                |  |||||  :  | ||||  :|         |:|| |:: ::| |||||||   |:   |
Db        384 IKPRIVGGTASVRGEWPWQVTLHTTSPTQRHLCGGSIIGNQWILTAAHCFYGVESPKILR 443

Qy         92 DPSGWMVQFGQLTSMPSFWSLQAYYTRYFVSNIYLSPRY-LGNSPYDIALVKLSAPVTYT 150
                 || : |  ::    ||: :|          | :  :| :  | |||||:||   | ||
Db        444 VYSGILNQ-SEIKEDTSFFGVQ---------EIIIHDQYKMAESGYDIALLKLETTVNYT 493

Qy        151 KHIQPICLQASTFEFENRTDCWVTGWGYIKEDEALPSPHTLQEVQVAIINNSMCNHLFLK 210
                :|||| :         ||||||||||  |  : :   :|||: :: :: |  |     :
Db        494 DSQRPICLPSKGDRNVIYTDCWVTGWGYRKLRDKI--QNTLQKAKIPLVTNEECQK---R 548

Qy        211 YSFRKDIFGDMVCAGNAQGGKDACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGV 270
                |  | |   |:|||  :|||||| |||||||:|  : :|: :|: ||| || :  ||||
Db        549 YRGHK-ITHKMICAGYREGGKDACKGDSGGPLSCKHNEVWHLVGITSWGEGCAQRERPGV 607

Qy        271 YTNISHHFEWI 281
                |||:  : :||
Db        608 YTNVVEYVDWI 618


RESULT 15
S00845
hepsin (EC 3.4.21.-) - human
C;Species: Homo sapiens (man)
C;Date: 31-Dec-1988 #sequence_revision 31-Dec-1988 #text_change 16-Aug-2004
C;Accession: S00845
R;Leytus, S.P.; Loeb, K.R.; Hagen, F.S.; Kurachi, K.; Davie, E.W.
Biochemistry 27, 1067-1074, 1988
A;Title: A novel trypsin-like serine protease (hepsin) with a putative transmembrane dom
A;Reference number: S00845; MUID:88209431; PMID:2835076
A;Accession: S00845
A;Molecule type: mRNA
A;Residues: 1-417 <LEY>
A;Cross-references: UNIPROT:P05981; EMBL:X07732; NID:g32063; PIDN:CAA30558.1; PID:g32064
C;Genetics:
A;Gene: GDB:HPN; TMPRSS1; hepsin
A;Cross-references: GDB:135685; OMIM:142440
A;Map position: 19q11-19q13.2
C;Superfamily: trypsin homology
C;Keywords: hydrolase; liver; serine proteinase; transmembrane protein
F;23-45/Domain: transmembrane #status predicted <TMN>
F;163-400/Domain: trypsin homology <TRY>
F;188-204,291-359,322-338,349-381/Disulfide bonds: #status predicted
F;203,257,353/Active site: His, Asp, Ser #status predicted

  Query Match             27.6%;  Score 476.5;  DB 1;  Length 417;
  Best Local Similarity   41.1%;  Pred. No. 6.2e-35;
  Matches  115;  Conservative   30;  Mismatches   96;  Indels   39;  Gaps   11;

Qy         21 PESQEAAPLSGPCGRRVI-TSRIVGGEDAELGRWPWQGSLRLWDSHVCGVSLLSHRWALT 79
                |  :  | :   |||| :   ||||| |  ||||||| |||   :|:|| ||||  | ||
Db        141 PRGRFLAAICQDCGRRKLPVDRIVGGRDTSLGRWPWQVSLRYDGAHLCGGSLLSGDWVLT 200

Qy         80 AAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQ------AYYTRYFVSNIYLSPRYLGN 133
                |||||   : :   | | | |    :  |  ||       |:  |    :  |   |
Db        201 AAHCFPERNRVL--SRWRV-FAGAVAQASPHGLQLGVQAVVYHGGYLP---FRDPNSEEN 254

Qy        134 SPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWG---YIKEDEALPSPHT 190
                |  ||||| ||:|:  |::|||:|| |:     :   | |||||   |  :   :
Db        255 S-NDIALVHLSSPLPLTEYIQPVCLPAAGQALVDGKICTVTGWGNTQYYGQQAGV----- 308
```

```
Qy        191 LQEVQVAIINNSMCNHLFLKYSFRKDIFGD-----MVCAGNAQGGKDACFGDSGGPLAC- 244
              ||| :| ||:| :||          | :|:     | |||  :|| ||| ||||||  |
Db        309 LQEARVPIISNDVCN--------GADFYGNQIKPKMFCAGYPEGGIDACQGDSGGPFVCE 360

Qy        245 ---NKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWI 281
                 ::   |   |:|||| ||    :||||| :|   |||
Db        361 DSISRTPRWRLCGIVSWGTGCALAQKPGVYTKVSDFREWI 400


Search completed: February 27, 2005, 20:23:27
Job time : 26.5815 secs
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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: February 27, 2005, 20:22:41 ; Search time 89.2843 Seconds
(without alignments)
1153.685 Million cell updates/sec

Title: US-10-040-647-6
Perfect score: 1728
Sequence: 1 MGARGALLLALLLARAGLRK.....PSWPLLFFPLLWALPLLGPV 314

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1385339 seqs, 328044528 residues

Total number of hits satisfying chosen parameters: 1385339

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_AA:*
1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
5: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*
6: /cgn2_6/ptodata/1/pubpaa/PCTUS_PUBCOMB.pep:*
7: /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
8: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
9: /cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
10: /cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep:*
11: /cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
12: /cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep:*
13: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
14: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
15: /cgn2_6/ptodata/1/pubpaa/US10C_PUBCOMB.pep:*
16: /cgn2_6/ptodata/1/pubpaa/US10D_PUBCOMB.pep:*
17: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
18: /cgn2_6/ptodata/1/pubpaa/US11_NEW_PUB.pep:*
19: /cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep:*
20: /cgn2_6/ptodata/1/pubpaa/US60_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1728	100.0	314	14	US-10-040-647-6	Sequence 6, Appli
2	1723	99.7	314	9	US-09-968-415-3	Sequence 3, Appli
3	1723	99.7	314	9	US-09-909-320-257	Sequence 257, App
4	1723	99.7	314	9	US-09-885-441-14	Sequence 14, Appl
5	1723	99.7	314	9	US-09-909-088B-257	Sequence 257, App
6	1723	99.7	314	9	US-09-905-291A-257	Sequence 257, App
7	1723	99.7	314	9	US-09-902-853-257	Sequence 257, App
8	1723	99.7	314	9	US-09-907-824-257	Sequence 257, App
9	1723	99.7	314	9	US-09-907-841-257	Sequence 257, App
10	1723	99.7	314	10	US-09-904-011-257	Sequence 257, App
11	1723	99.7	314	10	US-09-903-640-257	Sequence 257, App
12	1723	99.7	314	10	US-09-908-093-257	Sequence 257, App
13	1723	99.7	314	10	US-09-906-742-257	Sequence 257, App
14	1723	99.7	314	10	US-09-906-838-257	Sequence 257, App
15	1723	99.7	314	10	US-09-907-613-257	Sequence 257, App
16	1723	99.7	314	10	US-09-907-942-257	Sequence 257, App
17	1723	99.7	314	10	US-09-904-859-257	Sequence 257, App
18	1723	99.7	314	10	US-09-909-204-257	Sequence 257, App
19	1723	99.7	314	10	US-09-904-820-257	Sequence 257, App
20	1723	99.7	314	10	US-09-904-786-257	Sequence 257, App
21	1723	99.7	314	10	US-09-906-646-257	Sequence 257, App
22	1723	99.7	314	10	US-09-906-700-257	Sequence 257, App
23	1723	99.7	314	10	US-09-903-786-257	Sequence 257, App
24	1723	99.7	314	10	US-09-902-903-257	Sequence 257, App
25	1723	99.7	314	10	US-09-903-749A-257	Sequence 257, App
26	1723	99.7	314	10	US-09-904-119-257	Sequence 257, App
27	1723	99.7	314	10	US-09-904-956-257	Sequence 257, App
28	1723	99.7	314	10	US-09-902-736-257	Sequence 257, App
29	1723	99.7	314	10	US-09-907-794-257	Sequence 257, App
30	1723	99.7	314	10	US-09-903-943-257	Sequence 257, App
31	1723	99.7	314	10	US-09-904-462-257	Sequence 257, App
32	1723	99.7	314	10	US-09-907-925-257	Sequence 257, App
33	1723	99.7	314	10	US-09-902-692-257	Sequence 257, App
34	1723	99.7	314	10	US-09-903-520-257	Sequence 257, App
35	1723	99.7	314	10	US-09-905-056-257	Sequence 257, App
36	1723	99.7	314	10	US-09-909-064-257	Sequence 257, App
37	1723	99.7	314	10	US-09-904-553-257	Sequence 257, App
38	1723	99.7	314	10	US-09-905-381-257	Sequence 257, App
39	1723	99.7	314	10	US-09-904-485-257	Sequence 257, App
40	1723	99.7	314	10	US-09-905-348-257	Sequence 257, App
41	1723	99.7	314	10	US-09-905-088-257	Sequence 257, App
42	1723	99.7	314	10	US-09-907-575-257	Sequence 257, App
43	1723	99.7	314	10	US-09-905-075-257	Sequence 257, App
44	1723	99.7	314	10	US-09-902-759-257	Sequence 257, App
45	1723	99.7	314	10	US-09-902-634-257	Sequence 257, App

ALIGNMENTS

RESULT 1
US-10-040-647-6
; Sequence 6, Application US/10040647
; Publication No. US20030092154A1
; GENERAL INFORMATION:
; APPLICANT: (US only) ANTALIS Toni Marie and HOOPER John David
; TITLE OF INVENTION: NOVEL MOLECULES
; NUMBER OF SEQUENCES: 30
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: SCULLY, SCOTT, MURPHY & PRESSER
; STREET: 400 GARDEN CITY PLAZA
; CITY: GARDEN CITY
; STATE: NEW YORK
; COUNTRY: USA
; ZIP: 11530
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/10/040,647
; FILING DATE: 07-Jan-2002
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 09/023,942
; FILING DATE: <Unknown>
; APPLICATION NUMBER: PO5101/97
; FILING DATE: 13-FEB-1997
; APPLICATION NUMBER: PP0422/97
; FILING DATE: 18-NOV-1997
; APPLICATION NUMBER: International PCT Application
; FILING DATE: 13-FEB-1998
; ATTORNEY/AGENT INFORMATION:
; NAME: DIGIGLIO, FRANK S
; REGISTRATION NUMBER: 31,346

```
;            REFERENCE/DOCKET NUMBER: 11168
;       TELECOMMUNICATION INFORMATION:
;            TELEPHONE: (516) 742 4343
;            TELEFAX: (516) 742 4366
;            TELEX: 230 901 SANS UR
;   INFORMATION FOR SEQ ID NO: 6:
;       SEQUENCE CHARACTERISTICS:
;            LENGTH: 314 amino acids
;            TYPE: amino acid
;            TOPOLOGY: linear
;       MOLECULE TYPE: protein
;       SEQUENCE DESCRIPTION: SEQ ID NO: 6:
US-10-040-647-6

  Query Match            100.0%;  Score 1728;  DB 14;  Length 314;
  Best Local Similarity  100.0%;  Pred. No. 3.7e-159;
  Matches  314;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
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Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
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Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
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Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
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Db        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314


RESULT 2
US-09-968-415-3
; Sequence 3, Application US/09968415
; Publication No. US20020086334A1
;   GENERAL INFORMATION:
;       APPLICANT: Bandman, Olga
;                  Hillman, Jennifer L.
;                  Yue, Henry
;                  Guegler, Karl J.
;                  Corley, Neil C.
;                  Tang, Tom Y.
;                  Shah, Purvi
;       TITLE OF INVENTION: HUMAN PROTEASE MOLECULES
;       NUMBER OF SEQUENCES: 24
;       CORRESPONDENCE ADDRESS:
;            ADDRESSEE: Incyte Pharmaceuticals, Inc.
;            STREET: 3174 Porter Dr.
;            CITY: Palo Alto
;            STATE: CA
;            COUNTRY: USA
;            ZIP: 94304
;       COMPUTER READABLE FORM:
;            MEDIUM TYPE: Diskette
;            COMPUTER: IBM Compatible
;            OPERATING SYSTEM: DOS
;            SOFTWARE: FastSEQ for Windows Version 2.0
;       CURRENT APPLICATION DATA:
;            APPLICATION NUMBER: US/09/968,415
;            FILING DATE: 26-Sep-2001
;       PRIOR APPLICATION DATA:
;            APPLICATION NUMBER: 09/659,151
;            FILING DATE: <Unknown>
;       ATTORNEY/AGENT INFORMATION:
;            NAME: Mohan-Peterson, Sheela
;            REGISTRATION NUMBER: 41,201
;            REFERENCE/DOCKET NUMBER: PF-0458 US
;       TELECOMMUNICATION INFORMATION:
;            TELEPHONE: 650-855-0555
;            TELEFAX: 650-845-4166
;   INFORMATION FOR SEQ ID NO: 3:
;       SEQUENCE CHARACTERISTICS:
;            LENGTH: 314 amino acids
;            TYPE: amino acid
;            STRANDEDNESS: single
;            TOPOLOGY: linear
;       IMMEDIATE SOURCE:
;            LIBRARY: PROSTUT03
;            CLONE: 789927
;       SEQUENCE DESCRIPTION: SEQ ID NO: 3 :
US-09-968-415-3

  Query Match            99.7%;  Score 1723;  DB 9;  Length 314;
  Best Local Similarity  99.7%;  Pred. No. 1.1e-158;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
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Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
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Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314


RESULT 3
US-09-909-320-257
; Sequence 257, Application US/09909320
; Patent No. US20020132240A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
```

; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/909,320
; CURRENT FILING DATE: 2002-01-04
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
; LENGTH: 314
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-909-320-257

Query Match 99.7%; Score 1723; DB 9; Length 314;
Best Local Similarity 99.7%; Pred. No. 1.1e-158;
Matches 313; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

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Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
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Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314
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RESULT 4
US-09-885-441-14
; Sequence 14, Application US/09885441
; Patent No. US20020146407A1
; GENERAL INFORMATION:
; APPLICANT: Xiao, Yonghong
; TITLE OF INVENTION: Regulation of Human Eosinophil Serine
; TITLE OF INVENTION: Protease-1-Like Enzyme
; FILE REFERENCE: 04974.00512
; CURRENT APPLICATION NUMBER: US/09/885,441
; CURRENT FILING DATE: 2001-06-21
; PRIOR APPLICATION NUMBER: US 60/212,844
; PRIOR FILING DATE: 2000-06-21
; PRIOR APPLICATION NUMBER: US 60/244,171
; PRIOR FILING DATE: 2000-10-31
; PRIOR APPLICATION NUMBER: US 60/279,766
; PRIOR FILING DATE: 2001-03-30
; PRIOR APPLICATION NUMBER: PCT/_____
; PRIOR FILING DATE: 2001-06-20
; NUMBER OF SEQ ID NOS: 58
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 14
; LENGTH: 314
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-885-441-14

Query Match 99.7%; Score 1723; DB 9; Length 314;
Best Local Similarity 99.7%; Pred. No. 1.1e-158;
Matches 313; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

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Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
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Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
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Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
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Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
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Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314
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RESULT 5
US-09-909-088B-257
; Sequence 257, Application US/09909088B
; Patent No. US20020146709A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David

; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/909,088B
; CURRENT FILING DATE: 2001-07-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
; LENGTH: 314
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-909-088B-257

Query Match 99.7%; Score 1723; DB 9; Length 314;
Best Local Similarity 99.7%; Pred. No. 1.1e-158;
Matches 313; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

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Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
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Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314
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RESULT 6
US-09-905-291A-257
; Sequence 257, Application US/09905291A
; Patent No. US20020160374A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/905,291A
; CURRENT FILING DATE: 2001-07-12
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15

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;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
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;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-905-291A-257

  Query Match             99.7%;  Score 1723;  DB 9;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1.1e-158;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314


RESULT 7
US-09-902-853-257
; Sequence 257, Application US/09902853
; Publication No. US20020192659A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/902,853
;  CURRENT FILING DATE:  2001-07-10
;  PRIOR APPLICATION NUMBER: US/09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-902-853-257

  Query Match             99.7%;  Score 1723;  DB 9;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1.1e-158;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:|||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314


RESULT 8
US-09-907-824-257
; Sequence 257, Application US/09907824
; Publication No. US20020197671A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/907,824
;  CURRENT FILING DATE:  2001-07-17
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-907-824-257

  Query Match             99.7%;  Score 1723;  DB 9;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1.1e-158;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:|||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314


RESULT 9
US-09-907-841-257
; Sequence 257, Application US/09907841
; Publication No. US20020198366A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
```

```
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/907,841
;  CURRENT FILING DATE:  2001-11-20
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  Remaining Prior Application data removed - See File Wrapper or PALM.
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-907-841-257

  Query Match             99.7%;  Score 1723;  DB 9;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1.1e-158;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314


RESULT 10
US-09-904-011-257
; Sequence 257, Application US/09904011
; Publication No. US20030003530A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/904,011
;  CURRENT FILING DATE:  2001-07-11
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-904-011-257
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Query Match 99.7%; Score 1723; DB 10; Length 314;
Best Local Similarity 99.7%; Pred. No. 1.1e-158;
Matches 313; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

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Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314
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RESULT 11
US-09-903-640-257
; Sequence 257, Application US/09903640
; Publication No. US20030017463A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/903,640
; CURRENT FILING DATE: 2001-07-11
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
; LENGTH: 314
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-903-640-257

Query Match 99.7%; Score 1723; DB 10; Length 314;
Best Local Similarity 99.7%; Pred. No. 1.1e-158;
Matches 313; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
Db 1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy 61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
Db 61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy 121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
Db 121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy 181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
Db 181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy 241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
Db 241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy 301 FFPLLWALPLLGPV 314
Db 301 FFPLLWALPLLGPV 314

RESULT 12
US-09-908-093-257
; Sequence 257, Application US/09908093
; Publication No. US20030017498A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/908,093
; CURRENT FILING DATE: 2001-07-17
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222

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;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-908-093-257

  Query Match           99.7%;  Score 1723;  DB 10;  Length 314;
  Best Local Similarity  99.7%;  Pred. No. 1.1e-158;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314


RESULT 13
US-09-906-742-257
; Sequence 257, Application US/09906742
; Publication No. US20030023054A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,742
;  CURRENT FILING DATE:  2001-07-16
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-906-742-257

  Query Match           99.7%;  Score 1723;  DB 10;  Length 314;
  Best Local Similarity  99.7%;  Pred. No. 1.1e-158;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
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                 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
                 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
                 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
                 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
                 |||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314


RESULT 14
US-09-906-838-257
; Sequence 257, Application US/09906838
; Publication No. US20030027143A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,838
;  CURRENT FILING DATE:  2001-07-16
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-906-838-257

  Query Match             99.7%;  Score 1723;  DB 10;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1.1e-158;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
                 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
                 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
                 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
                 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
                 |||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314


RESULT 15
US-09-907-613-257
; Sequence 257, Application US/09907613
; Publication No. US20030027145A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
```

; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/907,613
; CURRENT FILING DATE: 2001-07-17
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
; LENGTH: 314
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-907-613-257

Query Match 99.7%; Score 1723; DB 10; Length 314;
Best Local Similarity 99.7%; Pred. No. 1.1e-158;
Matches 313; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

```
Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314
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Job time : 90.2843 secs

GenCore version 5.1.6

OM protein - protein search, using sw model

Run on: February 27, 2005, 20:12:15 ; Search time 30.0958 Seconds
(without alignments)
778.839 Million cell updates/sec

Title: US-10-040-647-6
Perfect score: 1728
Sequence: 1 MGARGALLLALLLARAGLRK.....PSWPLLFFPLLWALPLLGPV 314

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 513545 seqs, 74649064 residues

Total number of hits satisfying chosen parameters: 513545

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_AA:*
1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
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6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1728	100.0	314	4	US-09-023-942A-6	Sequence 6, Appli
2	1723	99.7	314	3	US-09-008-271A-3	Sequence 3, Appli
3	1723	99.7	314	4	US-09-907-794A-257	Sequence 257, App
4	1723	99.7	314	4	US-09-905-125A-257	Sequence 257, App
5	1723	99.7	314	4	US-09-902-775A-257	Sequence 257, App
6	1723	99.7	314	4	US-09-906-700-257	Sequence 257, App
7	1723	99.7	314	4	US-09-903-603A-257	Sequence 257, App
8	1723	99.7	314	4	US-09-904-920A-257	Sequence 257, App
9	1723	99.7	314	4	US-09-909-064-257	Sequence 257, App
10	1723	99.7	314	4	US-09-905-381A-257	Sequence 257, App
11	1723	99.7	314	4	US-09-906-618-257	Sequence 257, App
12	1706	98.7	312	4	US-09-023-942A-4	Sequence 4, Appli
13	1414	81.8	306	4	US-09-386-642-53	Sequence 53, Appl
14	1113.5	64.4	285	4	US-09-023-942A-26	Sequence 26, Appl
15	573	33.2	290	4	US-09-386-653A-7	Sequence 7, Appli
16	556	32.2	299	3	US-08-944-483-66	Sequence 66, Appl
17	549	31.8	315	4	US-09-386-653A-9	Sequence 9, Appli
18	543.5	31.5	328	4	US-09-386-642-11	Sequence 11, Appl
19	542.5	31.4	319	4	US-09-386-642-12	Sequence 12, Appl
20	536.5	31.0	317	4	US-09-386-629-7	Sequence 7, Appli
21	536.5	31.0	317	4	US-09-907-794A-263	Sequence 263, App
22	536.5	31.0	317	4	US-09-905-125A-263	Sequence 263, App
23	536.5	31.0	317	4	US-09-902-775A-263	Sequence 263, App
24	536.5	31.0	317	4	US-09-906-700-263	Sequence 263, App
25	536.5	31.0	317	4	US-09-903-603A-263	Sequence 263, App
26	536.5	31.0	317	4	US-09-904-920A-263	Sequence 263, App
27	536.5	31.0	317	4	US-09-909-064-263	Sequence 263, App
28	536.5	31.0	317	4	US-09-905-381A-263	Sequence 263, App
29	536.5	31.0	317	4	US-09-906-618-263	Sequence 263, App
30	523.5	30.3	270	2	US-08-978-404B-8	Sequence 8, Appli
31	511.5	29.6	273	2	US-08-978-404B-6	Sequence 6, Appli
32	511	29.6	276	2	US-09-016-366A-15	Sequence 15, Appl
33	511	29.6	276	2	US-08-978-404B-21	Sequence 21, Appl
34	511	29.6	327	4	US-09-386-629-8	Sequence 8, Appli
35	510	29.5	273	2	US-09-016-366A-19	Sequence 19, Appl
36	510	29.5	273	2	US-08-978-404B-14	Sequence 14, Appl
37	510	29.5	638	2	US-08-681-151-3	Sequence 3, Appli
38	509	29.5	273	2	US-08-978-404B-3	Sequence 3, Appli
39	509	29.5	274	2	US-09-016-366A-21	Sequence 21, Appl
40	509	29.5	274	2	US-08-978-404B-16	Sequence 16, Appl
41	508.5	29.4	446	4	US-10-177-661-4	Sequence 4, Appli
42	508.5	29.4	477	4	US-10-177-661-2	Sequence 2, Appli
43	508.5	29.4	562	4	US-09-879-792-12	Sequence 12, Appl
44	507	29.3	267	2	US-09-016-366A-23	Sequence 23, Appl
45	507	29.3	267	2	US-08-978-404B-18	Sequence 18, Appl

ALIGNMENTS

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RESULT 1
US-09-023-942A-6
; Sequence 6, Application US/09023942A
; Patent No. 6479274
;  GENERAL INFORMATION:
;    APPLICANT:  (US only) ANTALIS Toni Marie and HOOPER John David
;    TITLE OF INVENTION:  NOVEL MOLECULES
;    NUMBER OF SEQUENCES:  30
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  SCULLY, SCOTT, MURPHY & PRESSER
;      STREET:  400 GARDEN CITY PLAZA
;      CITY:  GARDEN CITY
;      STATE:  NEW YORK
;      COUNTRY:  USA
;      ZIP:  11530
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/023,942A
;      FILING DATE:  13-FEB-1998
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PO5101/97
;      FILING DATE:  13-FEB-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PP0422/97
;      FILING DATE:  18-NOV-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  International PCT Application
;      FILING DATE:  13-FEB-1998
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  DIGIGLIO,  FRANK S
;      REGISTRATION NUMBER:  31,346
;      REFERENCE/DOCKET NUMBER:  11168
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (516) 742 4343
;      TELEFAX:  (516) 742 4366
;      TELEX:  230 901 SANS UR
;  INFORMATION FOR SEQ ID NO:  6:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  314 amino acids
;      TYPE:  amino acid
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  protein
US-09-023-942A-6

  Query Match             100.0%;  Score 1728;  DB 4;  Length 314;
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  Best Local Similarity   100.0%;  Pred. No. 1.1e-165;
  Matches  314;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

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Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
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Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314


RESULT 2
US-09-008-271A-3
; Sequence 3, Application US/09008271A
; Patent No. 6203979
;   GENERAL INFORMATION:
;        APPLICANT: Bandman, Olga
;                   Hillman, Jennifer L.
;                   Yue, Henry
;                   Guegler, Karl J.
;                   Corley, Neil C.
;                   Tang, Tom Y.
;                   Shah, Purvi
;        TITLE OF INVENTION: HUMAN PROTEASE MOLECULES
;        NUMBER OF SEQUENCES: 24
;        CORRESPONDENCE ADDRESS:
;             ADDRESSEE: Incyte Pharmaceuticals, Inc.
;             STREET: 3174 Porter Dr.
;             CITY: Palo Alto
;             STATE: CA
;             COUNTRY: USA
;             ZIP: 94304
;        COMPUTER READABLE FORM:
;             MEDIUM TYPE: Diskette
;             COMPUTER: IBM Compatible
;             OPERATING SYSTEM: DOS
;             SOFTWARE: FastSEQ for Windows Version 2.0
;        CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/09/008,271A
;             FILING DATE: 16-Jan-1998
;        PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: <Unknown>
;             FILING DATE: <Unknown>
;        ATTORNEY/AGENT INFORMATION:
;             NAME: Mohan-Peterson, Sheela
;             REGISTRATION NUMBER: 41,201
;             REFERENCE/DOCKET NUMBER: PF-0458 US
;        TELECOMMUNICATION INFORMATION:
;             TELEPHONE: 650-855-0555
;             TELEFAX: 650-845-4166
;   INFORMATION FOR SEQ ID NO: 3:
;        SEQUENCE CHARACTERISTICS:
;             LENGTH: 314 amino acids
;             TYPE: amino acid
;             STRANDEDNESS: single
;             TOPOLOGY: linear
;        IMMEDIATE SOURCE:
;             LIBRARY: PROSTUT03
;             CLONE: 789927
;        SEQUENCE DESCRIPTION: SEQ ID NO: 3 :
US-09-008-271A-3

  Query Match            99.7%;  Score 1723;  DB 3;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 3.5e-165;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
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Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
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Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314


RESULT 3
US-09-907-794A-257
; Sequence 257, Application US/09907794A
; Patent No. 6635468
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/907,794A
;  CURRENT FILING DATE:  2001-07-17
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
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;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-907-794A-257

  Query Match             99.7%;  Score 1723;  DB 4;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 3.5e-165;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
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Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
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Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
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Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314


RESULT 4
US-09-905-125A-257
; Sequence 257, Application US/09905125A
; Patent No. 6664376
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/905,125A
;  CURRENT FILING DATE:  2001-07-12
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-905-125A-257

  Query Match             99.7%;  Score 1723;  DB 4;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 3.5e-165;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;
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Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
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Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
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Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
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Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
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Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314
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RESULT 5
US-09-902-775A-257
; Sequence 257, Application US/09902775A
; Patent No. 6686451
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/902,775A
; CURRENT FILING DATE: 2001-07-10
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
; LENGTH: 314
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-902-775A-257

Query Match 99.7%; Score 1723; DB 4; Length 314;
Best Local Similarity 99.7%; Pred. No. 3.5e-165;
Matches 313; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

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Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
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Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
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Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
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Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
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Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314
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RESULT 6
US-09-906-700-257
; Sequence 257, Application US/09906700
; Patent No. 6723535
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter

```
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,700
;  CURRENT FILING DATE:  2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-906-700-257

  Query Match           99.7%;  Score 1723;  DB 4;  Length 314;
  Best Local Similarity  99.7%;  Pred. No. 3.5e-165;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
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Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
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Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
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Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
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Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
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Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314


RESULT 7
US-09-903-603A-257
; Sequence 257, Application US/09903603A
; Patent No. 6767995
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: GNE.1618P2C12
;  CURRENT APPLICATION NUMBER: US/09/903,603A
;  CURRENT FILING DATE:  2001-07-11
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
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;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-903-603A-257

  Query Match             99.7%;  Score 1723;  DB 4;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 3.5e-165;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              |||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314


RESULT 8
US-09-904-920A-257
; Sequence 257, Application US/09904920A
; Patent No. 6806352
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/904,920A
;  CURRENT FILING DATE:  2001-07-13
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-904-920A-257

  Query Match             99.7%;  Score 1723;  DB 4;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 3.5e-165;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
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Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
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Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
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Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
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Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
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Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314


RESULT 9
US-09-909-064-257
; Sequence 257, Application US/09909064
; Patent No. 6818449
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/909,064
;  CURRENT FILING DATE:  2001-07-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
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;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-909-064-257

  Query Match             99.7%;  Score 1723;  DB 4;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 3.5e-165;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314


RESULT 10
US-09-905-381A-257
; Sequence 257, Application US/09905381A
; Patent No. 6818746
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
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;  CURRENT APPLICATION NUMBER: US/09/905,381A
;  CURRENT FILING DATE:  2001-07-13
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-905-381A-257

  Query Match             99.7%;  Score 1723;  DB 4;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 3.5e-165;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              |||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314


RESULT 11
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US-09-906-618-257
; Sequence 257, Application US/09906618
; Patent No. 6828146
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,618
;  CURRENT FILING DATE:  2001-07-16
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 257
;   LENGTH: 314
;   TYPE: PRT
;   ORGANISM: Homo Sapien
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US-09-906-618-257

  Query Match             99.7%;  Score 1723;  DB 4;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 3.5e-165;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
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Db        301 FFPLLWALPLLGPV 314


RESULT 12
US-09-023-942A-4
; Sequence 4, Application US/09023942A
; Patent No. 6479274
;  GENERAL INFORMATION:
;    APPLICANT:  (US only) ANTALIS Toni Marie and HOOPER John David
;    TITLE OF INVENTION:  NOVEL MOLECULES
;    NUMBER OF SEQUENCES:  30
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  SCULLY, SCOTT, MURPHY & PRESSER
;      STREET:  400 GARDEN CITY PLAZA
;      CITY:  GARDEN CITY
;      STATE:  NEW YORK
;      COUNTRY:  USA
;      ZIP:  11530
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/023,942A
;      FILING DATE:  13-FEB-1998
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PO5101/97
;      FILING DATE:  13-FEB-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PP0422/97
;      FILING DATE:  18-NOV-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  International PCT Application
;      FILING DATE:  13-FEB-1998
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  DIGIGLIO,  FRANK S
;      REGISTRATION NUMBER:  31,346
;      REFERENCE/DOCKET NUMBER:  11168
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (516) 742 4343
;      TELEFAX:  (516) 742 4366
;      TELEX:  230 901 SANS UR
;  INFORMATION FOR SEQ ID NO:  4:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  312 amino acids
;      TYPE:  amino acid
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  protein
US-09-023-942A-4

  Query Match             98.7%;  Score 1706;  DB 4;  Length 312;
  Best Local Similarity   99.4%;  Pred. No. 1.8e-163;
  Matches  312;  Conservative    0;  Mismatches    0;  Indels    2;  Gaps    1;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              |||||||||||||||||||||||||  ||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFET--DLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 118

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        119 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 178

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        179 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 238

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        239 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 298

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        299 FFPLLWALPLLGPV 312


RESULT 13
US-09-386-642-53
; Sequence 53, Application US/09386642
; Patent No. 6420157
; GENERAL INFORMATION:
;  APPLICANT: Darrow, Andrew
;  APPLICANT: Qi, Jenson
;  APPLICANT: Andrade-Gordon, Patricia
;  TITLE OF INVENTION: Zymogen Activation System
;  FILE REFERENCE: ORT-1028
;  CURRENT APPLICATION NUMBER: US/09/386,642
;  CURRENT FILING DATE: 1999-08-31
;  NUMBER OF SEQ ID NOS: 60
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 53
;   LENGTH: 306
;   TYPE: PRT
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence: Fusion gene of
;   OTHER INFORMATION: human protease F in CFEK2 zymogen vector
US-09-386-642-53

  Query Match             81.8%;  Score 1414;  DB 4;  Length 306;
  Best Local Similarity   89.3%;  Pred. No. 4.4e-134;
  Matches  260;  Conservative    5;  Mismatches   20;  Indels    6;  Gaps    2;

Qy         10 ALLLARAGLRKP---ESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRLWDSHV 66
              |||    |   |   :  :|| |: |        :||||   ||||||||||||||||||
Db         11 ALLGTTFGCGVPDYKDDDDAAALAAPFDD---DDKIVGGYALELGRWPWQGSLRLWDSHV 67

Qy         67 CGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYFVSNIYL 126
              |||||||||||||||||||||||||||||||||||||||||||||||||| |||||||||
Db         68 CGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYNRYFVSNIYL 127

Qy        127 SPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIKEDEALP 186
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db        128 SPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIKEDEALP 187

Qy        187 SPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGGPLACNK 246
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        188 SPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGGPLACNK 247

Qy        247 DGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSW 297
              :|||||||||||||||||||||||||||||||||||||||||||||||||||
Db        248 NGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSW 298


RESULT 14
US-09-023-942A-26
; Sequence 26, Application US/09023942A
; Patent No. 6479274
;  GENERAL INFORMATION:
;    APPLICANT:  (US only) ANTALIS Toni Marie and HOOPER John David
;    TITLE OF INVENTION:  NOVEL MOLECULES
;    NUMBER OF SEQUENCES:  30
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  SCULLY, SCOTT, MURPHY & PRESSER
;      STREET:  400 GARDEN CITY PLAZA
;      CITY:  GARDEN CITY
;      STATE:  NEW YORK
;      COUNTRY:  USA
;      ZIP:  11530
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/023,942A
;      FILING DATE:  13-FEB-1998
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PO5101/97
;      FILING DATE:  13-FEB-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PP0422/97
;      FILING DATE:  18-NOV-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  International PCT Application
;      FILING DATE:  13-FEB-1998
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  DIGIGLIO,  FRANK S
;      REGISTRATION NUMBER:  31,346
;      REFERENCE/DOCKET NUMBER:  11168
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (516) 742 4343
;      TELEFAX:  (516) 742 4366
;      TELEX:  230 901 SANS UR
;  INFORMATION FOR SEQ ID NO:  26:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  285 amino acids
;      TYPE:  amino acid
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  protein
US-09-023-942A-26

  Query Match             64.4%;  Score 1113.5;  DB 4;  Length 285;
  Best Local Similarity   69.1%;  Pred. No. 7.3e-104;
  Matches  197;  Conservative   35;  Mismatches   50;  Indels    3;  Gaps    2;

Qy         29 LSGPCGRRVITSRIVGGEDAELGRWPWQGSLRLWDSHVCGVSLLSHRWALTAAHCFETYS 88
              |||||| | | |||||||:|||||||||||||||:| :|:|| :||: || |||||||:
Db          3 LSGPCGHRTIPSRIVGGDDAELGRWPWQGSLRVWGNHLCGATLLNRRWVLTAAHCFQ--K 60

Qy         89 DLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYFVSNIYLSPRYLGNSPYDIALVKLSAPVT 148
              | :||  | ||||:||| || |:||||  || : :|:|||:|    | ||||:|||:|||
Db         61 D-NDPFDWTVQFGELTSRPSLWNLQAYSNRYQIEDIFLSPKYSEQYPNDIALLKLSSPVT 119

Qy        149 YTKHIQPICLQASTFEFENRTDCWVTGWGYIKEDEALPSPHTLQEVQVAIINNSMCNHLF 208
              |  ||||||  ||::|||||||||||||| | |||:||||:|||||||||||||||::
Db        120 YNNFIQPICLLNSTYKFENRTDCWVTGWGAIGEDESLPSPNTLQEVQVAIINNSMCNHMY 179

Qy        209 LKYSFRKDIFGDMVCAGNAQGGKDACFGDSGGPLACNKDGLWYQIGVVSWGVGCGRPNRP 268
              |  || :|:|||||||  :|||||||||||||||::| :|||:||||||:||||||||
Db        180 KKPDFRTNIWGDMVCAGTPEGGKDACFGDSGGPLACDQDTVWYQVGVVSWGIGCGRPNRP 239

Qy        269 GVYTNISHHFEWIQKLMAQSGMSQPDPSWPLLFFPLLWALPLLGP 313
              |||||||||: ||| | ::|: :|||   |||  | ||  || |
Db        240 GVYTNISHHYNWIQSTMIRNGLLRPDPVPLLLFLTLAWASSLLRP 284


RESULT 15
US-09-386-653A-7
; Sequence 7, Application US/09386653A
; Patent No. 6458564
; GENERAL INFORMATION:
;  APPLICANT: Andrade-Gordon, Patricia
;  APPLICANT:  Darrow, Andrew
;  APPLICANT:  Qi, Jian-shen
;  TITLE OF INVENTION: DNA encoding the novel human serine
;  TITLE OF INVENTION:  protease T
;  FILE REFERENCE: ORT-1032
;  CURRENT APPLICATION NUMBER: US/09/386,653A
;  CURRENT FILING DATE:  1999-08-31
;  NUMBER OF SEQ ID NOS: 11
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 7
;   LENGTH: 290
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-386-653A-7

  Query Match             33.2%;  Score 573;  DB 4;  Length 290;
  Best Local Similarity   42.6%;  Pred. No. 2e-49;
  Matches  123;  Conservative   44;  Mismatches  100;  Indels   22;  Gaps    8;

Qy          4 RGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLRLWD 63
              | | :  |||   | :: ::  |       |||  : :|:|||:| : | |||| |::
Db          3 RPAAVPLLLLLCFGSQRAKAATA------CGRPRMLNRMVGGQDTQEGEWPWQVSIQRNG 56

Qy         64 SHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFG--QLTSMPSFWSLQAYYTRYFV 121
              || || ||:: :| |||||||    : |: | : |  |  ||          | | |  |
Db         57 SHFCGGSLIAEQWVLTAAHCFR---NTSETSLYQVLLGARQLVQP----GPHAMYAR--V 107

Qy        122 SNIYLSPRYLGN-SPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
               :  :| | |  |  |:|||:| ||| :| :| |:||   :  ||   :|||||||
Db        108 RQVESNPLYQGTASSADVALVELEAPVPFTNYILPVCLPDPSVIFETGMNCWVTGWGSPS 167

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLK---YSFR-KDIFGDMVCAGNAQGGKDACFG 236
              |:: || |  ||:: | ||:   || |: |   : :: | |  ||:|||  :| |||| |
Db        168 EEDLLPEPRILQKLAVPIIDTPKCNLLYSKDTEFGYQPKTIKNDMLCAGFEEGKKDACKG 227

Qy        237 DSGGPLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLM 285
              |||||| |      | | ||:||| || | ||||||  :: |  || :::
Db        228 DSGGPLVCLVGQSWLQAGVISWGEGCARQNRPGVYIRVTAHHNWIHRII 276


Search completed: February 27, 2005, 20:24:33
Job time : 31.0958 secs
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GenCore version 5.1.6

OM protein - protein search, using sw model

Run on: February 27, 2005, 20:03:40 ; Search time 117.374 Seconds
(without alignments)
1034.667 Million cell updates/sec

Title: US-10-040-647-6
Perfect score: 1728
Sequence: 1 MGARGALLLALLLARAGLRK.....PSWPLLFFPLLWALPLLGPV 314

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 2105692 seqs, 386760381 residues

Total number of hits satisfying chosen parameters: 2105692

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : A_Geneseq_16Dec04:*
1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
4: geneseqp2001s:*
5: geneseqp2002s:*
6: geneseqp2003as:*
7: geneseqp2003bs:*
8: geneseqp2004s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1728	100.0	314	2	AAW77297	Aaw77297 Amino aci
2	1723	99.7	314	2	AAW77296	Aaw77296 Amino aci
3	1723	99.7	314	2	AAW97116	Aaw97116 A human e
4	1723	99.7	314	2	AAY06434	Aay06434 Human pro
5	1723	99.7	314	2	AAY13388	Aay13388 Amino aci
6	1723	99.7	314	3	AAB12132	Aab12132 Hydrophob
7	1723	99.7	314	3	ADC78569	Adc78569 Human PRO
8	1723	99.7	314	4	AAB80256	Aab80256 Human PRO
9	1723	99.7	314	4	AAU02223	Aau02223 Human ext
10	1723	99.7	314	5	AAE17010	Aae17010 Human eos
11	1723	99.7	314	6	ABU71634	Abu71634 Human PRO
12	1723	99.7	314	6	ABU71489	Abu71489 Human PRO
13	1723	99.7	314	6	ABU71935	Abu71935 Human sec
14	1723	99.7	314	6	ABO01818	Abo01818 Novel hum
15	1723	99.7	314	6	ABU54391	Abu54391 Human sec
16	1723	99.7	314	6	ABO47406	Abo47406 Human sec
17	1723	99.7	314	6	ABU64543	Abu64543 Human sec
18	1723	99.7	314	6	ABU67389	Abu67389 Human sec
19	1723	99.7	314	6	ABO14909	Abo14909 Human sec
20	1723	99.7	314	6	ABU69666	Abu69666 Novel hum
21	1723	99.7	314	6	ABO14848	Abo14848 Human sec
22	1723	99.7	314	6	ADB29462	Adb29462 Human sec
23	1723	99.7	314	6	ADA18318	Ada18318 Human sec
24	1723	99.7	314	6	ABO32800	Abo32800 Human sec
25	1723	99.7	314	6	ABO34860	Abo34860 Human PRO
26	1723	99.7	314	6	ADA16293	Ada16293 Human sec
27	1723	99.7	314	6	ADA42438	Ada42438 Human sec
28	1723	99.7	314	6	ABO17538	Abo17538 Human PRO
29	1723	99.7	314	7	ADA16717	Ada16717 Human sec
30	1723	99.7	314	7	ADA13146	Ada13146 Human sec
31	1723	99.7	314	7	ADA42014	Ada42014 Human sec
32	1723	99.7	314	7	ADA17361	Ada17361 Human sec
33	1723	99.7	314	7	ADA42864	Ada42864 Human sec
34	1723	99.7	314	7	ABO17599	Abo17599 Human PRO
35	1723	99.7	314	7	ADB80572	Adb80572 Ovarian c
36	1723	99.7	314	7	ADB77783	Adb77783 Human sec
37	1723	99.7	314	7	ADB74919	Adb74919 Human sec
38	1723	99.7	314	7	ADC28565	Adc28565 Human sec
39	1723	99.7	314	7	ADC39765	Adc39765 Human sec
40	1723	99.7	314	7	ADC40279	Adc40279 Human sec
41	1723	99.7	314	7	ADC19103	Adc19103 Human sec
42	1723	99.7	314	7	ADC34403	Adc34403 Human sec
43	1723	99.7	314	7	ADC29458	Adc29458 Human sec
44	1723	99.7	314	7	ADC28989	Adc28989 Human sec
45	1723	99.7	314	7	ADC40874	Adc40874 Human sec

ALIGNMENTS

RESULT 1
AAW77297
ID AAW77297 standard; protein; 314 AA.
XX
AC AAW77297;
XX
DT 07-JAN-1999 (first entry)
XX
DE Amino acid sequence of long isoform of HELA2.
XX
KW Serine protease; regulation; cell activity; viability; HELA2; ATC2;
KW BCOM3; testisin; fertility; suppressor; testicular germ cell cancer;
KW seminoma; testis-specific expression; antitumour; sperm development;
KW infertility.
XX
OS Homo sapiens.
XX
PN WO9836054-A1.
XX
PD 20-AUG-1998.
XX
PF 13-FEB-1998; 98WO-AU000085.
XX
PR 13-FEB-1997; 97AU-00005101.
PR 18-NOV-1997; 97AU-00000422.
XX
PA (AMRA-) AMRAD OPERATIONS PTY LTD.
XX
PI Antalis TM, Hooper JD;
XX
DR WPI; 1998-480768/41.
DR N-PSDB; AAV59119.
XX
PT New serine protease(s) and kinase involved in regulating cell activity
PT and viability - particularly the testis-specific protease HELA2 used for
PT modulation of fertility and as tumour suppressor.
XX
PS Claim 3; Page 62-64; 167pp; English.
XX
CC The present sequence represents the amino acid sequence of the long
CC isoform of HELA2. cDNA generated from HeLa cells and PAI-2 expressing
CC HeLa cells was amplified using PCR primers AAV48312-13. Three new
CC sequences were detected in the 480 bp amplicon. These sequences are
CC designated HELA2 and ATC2 which have high homology to serine proteases
CC and BCOM3 which has homology to a kinase. The proteins are involved in or
CC associated with regulation of cell activity and/or viability.
CC Administration of recombinant HELA2 (also called testisin) is used to
CC increase fertility. Downregulation of HELA2 reduces fertility. HELA2 is

```
CC   also a suppressor of testicular germ cell cancers (seminoma) and is also
CC   expressed in some non-testicular cancers (of colon, pancreas, prostate
CC   and ovary), so is a marker/potential therapeutic target for cancer. The
CC   promoter from the HELA2 gene is useful for testis-specific expression of
CC   other genes, e.g. for gene therapy or modulation of fertility. Drugs that
CC   block activity of HELA2 should have antitumour activity (other than in
CC   testis) while in testis recombinant HELA2 should stop growth of tumours
CC   and normalise sperm development (eliminating the need for orchidectomy).
CC   Identification of mutant forms of HELA2 can be used to diagnose
CC   infertility
XX
SQ   Sequence 314 AA;

  Query Match             100.0%;  Score 1728;  DB 2;  Length 314;
  Best Local Similarity   100.0%;  Pred. No. 5.9e-147;
  Matches  314;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314


RESULT 2
AAW77296
ID   AAW77296 standard; protein; 314 AA.
XX
AC   AAW77296;
XX
DT   07-JAN-1999  (first entry)
XX
DE   Amino acid sequence of the short isoform of HELA2.
XX
KW   Serine protease; regulation; cell activity; viability; HELA2; ATC2;
KW   BCOM3; testisin; fertility; suppressor; testicular germ cell cancer;
KW   seminoma; testis-specific expression; antitumour; sperm development;
KW   infertility.
XX
OS   Homo sapiens.
XX
PN   WO9836054-A1.
XX
PD   20-AUG-1998.
XX
PF   13-FEB-1998;   98WO-AU000085.
XX
PR   13-FEB-1997;   97AU-00005101.
PR   18-NOV-1997;   97AU-00000422.
XX
PA   (AMRA-) AMRAD OPERATIONS PTY LTD.
XX
PI   Antalis TM,  Hooper JD;
XX
DR   WPI; 1998-480768/41.
DR   N-PSDB; AAV59118.
XX
PT   New serine protease(s) and kinase involved in regulating cell activity
PT   and viability - particularly the testis-specific protease HELA2 used for
PT   modulation of fertility and as tumour suppressor.
XX
PS   Claim 2; Fig 6; 167pp; English.
XX
CC   The present sequence represents the amino acid sequence of the short
CC   isoform of HELA2. cDNA generated from HeLa cells and PAI-2 expressing
CC   HeLa cells was amplified using PCR primers AAV48312-13. Three new
CC   sequences were detected in the 480 bp amplicon. These sequences are
CC   designated HELA2 and ATC2 which have high homology to serine proteases
CC   and BCOM3 which has homology to a kinase. The proteins are involved in or
CC   associated with regulation of cell activity and/or viability.
CC   Administration of recombinant HELA2 (also called testisin) is used to
CC   increase fertility. Downregulation of HELA2 reduces fertility. HELA2 is
CC   also a suppressor of testicular germ cell cancers (seminoma) and is also
CC   expressed in some non-testicular cancers (of colon, pancreas, prostate
CC   and ovary), so is a marker/potential therapeutic target for cancer. The
CC   promoter from the HELA2 gene is useful for testis-specific expression of
CC   other genes, e.g. for gene therapy or modulation of fertility. Drugs that
CC   block activity of HELA2 should have antitumour activity (other than in
CC   testis) while in testis recombinant HELA2 should stop growth of tumours
CC   and normalise sperm development (eliminating the need for orchidectomy).
CC   Identification of mutant forms of HELA2 can be used to diagnose
CC   infertility
XX
SQ   Sequence 314 AA;

  Query Match             99.7%;  Score 1723;  DB 2;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1.7e-146;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              |||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314


RESULT 3
AAW97116
ID   AAW97116 standard; protein; 314 AA.
XX
AC   AAW97116;
XX
DT   04-MAY-1999  (first entry)
XX
DE   A human eosinophil serine protease.
XX
KW   Human; eosinophil; serine protease; allergic disease; infectious disease;
KW   tumour; granulomatous disease; collagen disease; vascular inflammation.
XX
OS   Homo sapiens.
```

XX
PN JP11032768-A.
XX
PD 09-FEB-1999.
XX
PF 16-JUL-1997; 97JP-00191319.
XX
PR 16-JUL-1997; 97JP-00191319.
XX
PA (ONOY) ONO PHARM CO LTD.
XX
DR WPI; 1999-183825/16.
DR N-PSDB; AAX15336.
XX
PT New eosinophils serine protease - useful for prevention and treatment of
PT allergic, infectious, tumor, granulomatous and collagen diseases.
XX
PS Claim 1; Page 9-10; 18pp; Japanese.
XX
CC The present sequence represents a human eosinophil serine protease. The
CC protease is useful in drug compositions for the prevention and treatment
CC of allergic diseases, infectious diseases, tumour diseases, granulomatous
CC diseases, collagen diseases and vascular inflammation
XX
SQ Sequence 314 AA;

Query Match 99.7%; Score 1723; DB 2; Length 314;
Best Local Similarity 99.7%; Pred. No. 1.7e-146;
Matches 313; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
Db 1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy 61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
Db 61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy 121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
Db 121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy 181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
Db 181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy 241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
Db 241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy 301 FFPLLWALPLLGPV 314
Db 301 FFPLLWALPLLGPV 314

RESULT 4
AAY06434
ID AAY06434 standard; protein; 314 AA.
XX
AC AAY06434;
XX
DT 27-SEP-1999 (first entry)
XX
DE Human protease HUPM-3.
XX
KW Protease; human; HUPM-3; cell proliferation; cancer; immune disorder;
KW inflammation; therapy.
XX
OS Homo sapiens.
XX
FH Key Location/Qualifiers
FT Peptide 1. .19
FT /note= "putative signal peptide"
FT Protein 20. .314
FT /note= "putative mature protein"
FT Modified-site 39
FT /note= "protein kinase C phosphorylation site"
FT Modified-site 58
FT /note= "protein kinase C phosphorylation site"
FT Modified-site 73
FT /note= "protein kinase C phosphorylation site"
FT Active-site 82
FT Modified-site 86
FT /note= "casein kinase II phosphorylation site"
FT Modified-site 127
FT /note= "protein kinase C phosphorylation site"
FT Modified-site 134
FT /note= "casein kinase II phosphorylation site"
FT Modified-site 161
FT /note= "casein kinase II phosphorylation site"
FT Modified-site 167
FT /note= "N-glycosylated"
FT Modified-site 190
FT /note= "casein kinase II phosphorylation site"
FT Modified-site 200
FT /note= "N-glycosylated"
FT Modified-site 212
FT /note= "protein kinase C phosphorylation site"
FT Active-site 238
FT Modified-site 273
FT /note= "N-glycosylated"
FT Modified-site 291
FT /note= "casein kinase II phosphorylation site"
XX
PN WO9936550-A2.
XX
PD 22-JUL-1999.
XX
PF 12-JAN-1999; 99WO-US000655.
XX
PR 16-JAN-1998; 98US-00008271.
XX
PA (INCY-) INCYTE PHARM INC.
XX
PI Bandman O, Hillman JL, Yue H, Guegler KJ, Corley NC, Tang YT;
PI Shah P;
XX
DR WPI; 1999-430616/36.
DR N-PSDB; AAX87151.
XX
PT Novel human protease molecules useful in the treatment of developmental
PT disorders and/or cancers.
XX
PS Claim 1; Page 71-72; 90pp; English.
XX
CC The present sequence represents novel human protease HUPM-3, as deduced
CC from the consensus sequence (see AAX87151) of overlapping cDNA clones
CC obtained from various libraries. Northern analysis shows expression of
CC HUPM-3 in cardiovascular, haematopoietic and male reproductive cDNA
CC libraries. Approximately 86% of these libraries are associated with
CC neoplastic disorders. The invention provides 12 new human proteases, i.e.
CC HUPM-1 to -12 (see AAY06432-43), and the polynucleotides encoding them
CC (see AAX87149-60). Also provided are vectors, host cells and methods for
CC producing HUPM polypeptides, as well as agonists and antagonists of HUPM.
CC Methods for treating or preventing cell proliferative disorders and
CC immune disorders using HUPM or HUPM antagonists are claimed
XX
SQ Sequence 314 AA;

Query Match 99.7%; Score 1723; DB 2; Length 314;
Best Local Similarity 99.7%; Pred. No. 1.7e-146;
Matches 313; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
Db 1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

```
Qy          61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy         121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy         181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy         241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
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Db         241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy         301 FFPLLWALPLLGPV 314
               ||||||||||||||
Db         301 FFPLLWALPLLGPV 314
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RESULT 5
AAY13388
ID AAY13388 standard; protein; 314 AA.
XX
AC AAY13388;
XX
DT 25-JUN-1999 (first entry)
XX
DE Amino acid sequence of protein PRO303.
XX
KW Secreted protein; transmembrane protein; human; enterocolitis;
KW Zollinger-Ellison syndrome; gastrointestinal ulceration;
KW congenital microvillus atrophy; skin disease; cell growth;
KW abnormal keratinocyte differentiation; psoriasis; epithelial cancer;
KW Parkinson's disease; Alzheimer's disease; ALS; neuropathy; fibromodulin;
KW dermal scarring; Usher Syndrome; Atrophia areata; anti-thrombotic;
KW wound healing; tissue repair.
XX
OS Homo sapiens.
XX
PN WO9914328-A2.
XX
PD 25-MAR-1999.
XX
PF 16-SEP-1998; 98WO-US019330.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
XX
PA (GETH) GENENTECH INC.
XX
PI Wood WI, Gurney AL, Goddard A, Pennica D, Chen J, Yuan J;
XX
DR WPI; 1999-229533/19.
DR N-PSDB; AAX52259.
XX
PT New isolated human genes and polypeptides used in, e.g. treatment of
PT gastrointestinal ulceration.
XX
PS Claim 12; Fig 92; 320pp; English.
XX
CC AAY13344-403 represent secreted and transmembrane human proteins. The
CC cDNA sequences are obtained from cDNA libraries, prepared from fetal
CC lung, fetal kidney, fetal brain, fetal liver and fetal retina. The
CC encoded polypeptides have specific uses based on their homology to known
CC polypeptides, e.g. PRO211 and PRO217 can be used for disorders associated
CC with the preservation and maintenance of gastrointestinal mucosa and the
CC repair of acute and chronic mucosal lesions (e.g. enterocolitis,
CC Zollinger-Ellison syndrome, gastrointestinal ulceration and congenital
CC microvillus atrophy), skin diseases associated with abnormal keratinocyte
CC differentiation (e.g. psoriasis, epithelial cancers such as lung squamous
CC cell carcinoma of the vulva and gliomas), potent effects on cell growth
CC and development, diseases related to growth or survival of nerve cells
CC including Parkinson's disease, Alzheimer's disease, ALS, neuropathies or
CC cancer. PRO265 can be used as for fibromodulin, e.g. for reducing dermal
CC scarring. PRO264 can be used as a target for anti-tumor drugs. PRO533 may
CC be used in the treatment of Usher Syndrome or Atrophia areata; PRO269 can
CC be used as an anti-thrombotic agent; PRO287 polypeptides and portions may
CC have therapeutic applications in wound healing and tissue repair; PRO317
CC can be used for treating problems of the kidney, uterus, endometrium,
CC blood vessels, or related tissue, e.g. in the heart of genital tract
XX
SQ Sequence 314 AA;

```
  Query Match             99.7%;  Score 1723;  DB 2;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1.7e-146;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy           1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db           1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy          61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
```

```
Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314
```

RESULT 6
AAB12132
ID AAB12132 standard; protein; 314 AA.
XX
AC AAB12132;
XX
DT 02-FEB-2001 (first entry)
XX
DE Hydrophobic domain protein from clone HP03116 isolated from KB cells.
XX
KW Human; secreted protein; membrane protein; hydrophobic domain;
KW proliferation control; differentiation induction; material transport;
KW biophylaxis; signal receptor; ion channel; transporter; immunostimulant;
KW immunosuppressant; haematopoiesis regulator; chemotactic; chemokinetic;
KW haemostatic; thrombolytic; anti-inflammatory; tumour inhibition;
KW autoimmune disease; Alzheimer's disease; Parkinson's disease; cancer.
XX
OS Homo sapiens.
XX
PN WO200029448-A2.
XX
PD 25-MAY-2000.
XX
PF 17-NOV-1999; 99WO-JP006412.
XX
PR 17-NOV-1998; 98JP-00326255.
PR 22-DEC-1998; 98JP-00364315.
PR 16-MAR-1999; 99JP-00069811.
PR 27-APR-1999; 99JP-00119299.
PR 19-MAY-1999; 99JP-00138169.
XX
PA (SAGA) SAGAMI CHEM RES CENT.
PA (PROT-) PROTEGENE INC.
XX
PI Kato S, Kimura T;
XX
DR WPI; 2000-387753/33.
DR N-PSDB; AAA62005, AAA62015.
XX
PT Proteins comprising hydrophobic regions, such as secretory and membrane
PT proteins, useful in research and diagnostics and having various
PT activities e.g. immunomodulatory, antiinflammatory, chemokinetic,
PT hemostatic, thrombolytic.
XX
PS Claim 1; Page 238-240; 410pp; English.
XX
CC Secretory proteins play important roles in the proliferation control, the
CC differentiation induction, the material transport and the biophylaxis of
CC cells. Membrane proteins have important roles as signal receptors, ion
CC channels and transporters. The present sequence is a human protein which
CC has at least one hydrophobic domain. This protein may be a secretory or a
CC membrane protein. The present protein may have cytokine and cell
CC proliferation/differentiation activity, immune stimulating or suppressing
CC activity, haematopoiesis activity, tissue growth activity,
CC activin/inhibin activity, chemotactic/chemokinetic activity, haemostatic
CC and thrombolytic activity, anti-inflammatory activity and tumour
CC inhibition activity. The present protein could therefore be used for
CC treatment of autoimmune disease, Alzheimer's disease, Parkinson's
CC disease, and cancer
XX
SQ Sequence 314 AA;

```
  Query Match             99.7%;  Score 1723;  DB 3;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1.7e-146;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314
```

RESULT 7
ADC78569
ID ADC78569 standard; protein; 314 AA.
XX
AC ADC78569;
XX
DT 01-JAN-2004 (first entry)
XX
DE Human PRO303 protein.
XX
KW antiinflamatory; antiulcer; cytostatic; antipsoriatic; antiparkinsonian;
KW nootropic; neuroprotective; vasotropic; chemotaxic; angiogenic;
KW neurotrophic; osteopathic; antiasthmatic; antiarthritic; antirheumatic;
KW antiarteriosclerotic; cardiant; antidiabetic; cerebroprotective;
KW thrombolytic; immunomodulator; enterocolitis; Zollinger-Ellison syndrome;
KW gastrointestinal ulceration; psoriasis; cancer; Parkinson's disease;
KW Alzheimer's; ALS; neuropathy; dermal scarring; wound healing;
KW nerve repair; thrombosis; bone; cartilage formation; angiogenesis;
KW asthma; rheumatoid arthritis; multiple sclerosis; inflammatory disorder;
KW atherosclerosis; cardiac injury; infertility; premature aging; AIDS;
KW diabetes; stroke; gene therapy; transgenic; PRO; human.
XX
OS Homo sapiens.
XX
PN WO200015796-A2.
XX
PD 23-MAR-2000.
XX
PF 15-SEP-1999; 99WO-US021090.
XX
PR 16-SEP-1998; 98WO-US019330.
XX
PA (GETH) GENENTECH INC.
XX
PI Chen J, Goddard A, Gurney AL, Hillan K, Pennica D, Wood WI;
PI Yuan J;
XX
DR WPI; 2000-271434/23.

```
DR   N-PSDB; ADC78568.
XX
PT   Novel nucleic acids encoding secreted and transmembrane polypeptides with
PT   homology, e.g. to growth and cancer-associated antigens.
XX
PS   Claim 12; SEQ ID NO 257; 355pp; English.
XX
CC   The invention relates to a novel nucleic acid encoding a PRO polypeptide.
CC   The polypeptides and polynucleotides of the invention may be useful as
CC   research tools and as therapeutics for treating enterocolitis, Zollinger-
CC   Ellison syndrome, gastrointestinal ulceration, psoriasis, cancer,
CC   Parkinson's disease, Alzheimer's disease, ALS, neuropathies, dermal
CC   scarring and wound healing, nerve repair, thrombosis, bone and/or
CC   cartilage formation, angiogenesis, asthma, rheumatoid arthritis, multiple
CC   sclerosis, inflammatory disorders, atherosclerosis, cardiac injury,
CC   infertility, premature aging, AIDS, diabetes complications and stroke.
CC   The molecules may also be utilised during gene therapy procedures and
CC   transgenic animal production. The current sequence is that of the human
CC   PRO protein of the invention.
XX
SQ   Sequence 314 AA;

  Query Match             99.7%;  Score 1723;  DB 3;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1.7e-146;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314


RESULT 8
AAB80256
ID   AAB80256 standard; protein; 314 AA.
XX
AC   AAB80256;
XX
DT   24-APR-2001  (first entry)
XX
DE   Human PRO303 protein.
XX
KW   Human; PRO; dermatological; antipsoriatic; cytostatic; antiinflammatory;
KW   antiparkinsonian nootropic; neuroprotective; vulnerary; cardiant;
KW   antiangiogenic; vasotropic; antiasthmatic; antirheumatic; cancer;
KW   antiarthritic; antiinfertility; antidiabetic; antiviral; diabetes;
KW   ophthalmological; gene therapy; skin disease; gastrointestinal disorder;
KW   ischaemia; inflammation.
XX
OS   Homo sapiens.
XX
PN   WO200104311-A1.
XX
PD   18-JAN-2001.
XX
PF   22-FEB-2000; 2000WO-US004414.
XX
PR   07-JUL-1999;   99US-0143048P.
PR   26-JUL-1999;   99US-0145698P.
PR   28-JUL-1999;   99US-0146222P.
PR   08-SEP-1999;   99WO-US020594.
PR   13-SEP-1999;   99WO-US020944.
PR   15-SEP-1999;   99WO-US021090.
PR   15-SEP-1999;   99WO-US021547.
PR   05-OCT-1999;   99WO-US023089.
PR   29-NOV-1999;   99WO-US028214.
PR   30-NOV-1999;   99WO-US028313.
PR   02-DEC-1999;   99WO-US028564.
PR   02-DEC-1999;   99WO-US028565.
PR   16-DEC-1999;   99WO-US030095.
PR   20-DEC-1999;   99WO-US030911.
PR   20-DEC-1999;   99WO-US030999.
PR   05-JAN-2000; 2000WO-US000219.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Ashkenazi AJ,  Botstein D,  Desnoyers L,  Eaton DL,  Ferrara N;
PI   Filvaroff E,  Fong S,  Gao W,  Gerber H,  Gerritsen ME,  Goddard A;
PI   Godowski PJ,  Grimaldi CJ,  Gurney AL,  Hillan KJ,  Kljavin IJ;
PI   Mather JP,  Pan J,  Paoni NF,  Roy MA,  Stewart TA,  Tumas D;
PI   Williams PM,  Wood WI;
XX
DR   WPI; 2001-081051/09.
DR   N-PSDB; AAF72417.
XX
PT   Sixty one nucleic acids encoding PRO polypeptides which are useful in the
PT   treatment of skin diseases (e.g. psoriasis), cancers (e.g. lung squamous
PT   cell carcinoma) and neurodegenerative diseases (e.g. Alzheimer's
PT   disease).
XX
PS   Claim 1; Fig 92; 393pp; English.
XX
CC   The present sequence is one of sixty one novel secreted and transmembrane
CC   PRO polypeptides. The PRO polypeptides are useful for treating skin
CC   diseases (e.g. psoriasis), cancers (e.g. lung squamous cell carcinoma),
CC   gastrointestinal disorders (e.g. enterocolitis), neurodegenerative
CC   diseases (e.g. Alzheimer's disease, Parkinson's disease), wound repair,
CC   cardiovascular disorders (e.g. endometrial bleeding angiogenesis,
CC   ischaemias such as coronary ischaemia, atherosclerosis), inflammatory
CC   disorders (e.g. asthma, rheumatoid arthritis, multiple sclerosis),
CC   infertility, AIDS and diabetes and retinal disorders such as retinitis
CC   pigmentosum. The PRO nucleic acids have applications in molecular
CC   biology, including use as hybridization probes, and in chromosome and
CC   gene mapping
XX
SQ   Sequence 314 AA;

  Query Match             99.7%;  Score 1723;  DB 4;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1:7e-146;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
```

```
Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314
```

RESULT 9
AAU02223
ID AAU02223 standard; protein; 314 AA.
XX
AC AAU02223;
XX
DT 29-AUG-2001 (first entry)
XX
DE Human extracellular serine protease TADG-16.
XX
KW Human; extracellular serine protease; tumour antigen derived gene-16;
KW TADG-16; ovarian carcinoma; breast cancer; lung cancer; colon cancer;
KW prostate cancer; HLA type.
XX
OS Homo sapiens.
XX
FH Key Location/Qualifiers
FT Peptide 1. .17
FT /label= Signal_secretion_sequence
FT Protein 18. .314
FT /label= Secreted_TADG-16_protein
FT Region 76. .83
FT /note= "Region containing His residue of conserved
FT catalytic triad found in serine proteases"
FT Region 137. .141
FT /note= "Region containing Asp residue of conserved
FT catalytic triad found in serine proteases"
FT Region 236. .241
FT /note= "Region containing Ser residue of conserved
FT catalytic triad found in serine proteases"
XX
PN WO200127257-A1.
XX
PD 19-APR-2001.
XX
PF 13-OCT-2000; 2000WO-US028558.
XX
PR 14-OCT-1999; 99US-00418527.
XX
PA (UYAR-) UNIV ARKANSAS.
XX
PI O'brien TJ, Underwood LJ, Shigemasa K;
XX
DR WPI; 2001-273769/28.
DR N-PSDB; AAS01698.
XX
PT New tumor antigen-derived gene-16 protein, useful for diagnosis and
PT treatment of ovarian, breast, lung, colon and prostate cancer.
XX
PS Claim 3; Fig 4; 124pp; English.
XX
CC The present sequence represents a novel human extracellular serine
CC protease, tumour antigen derived gene-16 (TADG-16) protein. TADG-16 is
CC expressed in normal ovaries and testes and in certain ovarian carcinomas.
CC TADG-16 contains the conserved catalytic triad, His-Asp-Ser, and a signal
CC secretion sequence characteristic of the serine protease family. An
CC antisense oligonucleotide having a complementary sequence to the TADG-16
CC nucleic acid is useful for treating various cancers, including ovarian,
CC breast, lung, colon and prostate. The TADG-16 nucleic acid, TADG-16
CC protein and antibodies specific to TADG-16 are useful for the diagnosis
CC of cancer. TADG-16 protein or its fragments are useful for vaccinating an
CC individual against TADG-16. Numerous TADG-16 peptides (AAU02225-AAU02384)
CC are tested for their binding affinity to the 8 haplotypes HLA A0201, HLA
CC A0205, HLA A1, HLA A24, HLA B7, HLA B8, HLA B2702, and HLA B4403
XX
SQ Sequence 314 AA;

Query Match 99.7%; Score 1723; DB 4; Length 314;
Best Local Similarity 99.7%; Pred. No. 1.7e-146;
Matches 313; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

```
Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314
```

RESULT 10
AAE17010
ID AAE17010 standard; protein; 314 AA.
XX
AC AAE17010;
XX
DT 18-APR-2002 (first entry)
XX
DE Human eosinophil serine protease-1 (esp-1) like enzyme #2.
XX
KW Human; eosinophil serine protease-1; esp-1; enzyme; antiinflammatory;
KW antiasthmatic; antiallergic; osteopathic; cytostatic; dermatological;
KW asthma; airway allergy; chronic obstructive pulmonary disease; COPD;
KW osteoporosis; dermatitis; Paget's disease; therapy.
XX
OS Homo sapiens.
XX
PN WO200198503-A2.
XX
PD 27-DEC-2001.
XX
PF 20-JUN-2001; 2001WO-EP006936.
XX
PR 21-JUN-2000; 2000US-0212844P.
PR 31-OCT-2000; 2000US-0244171P.
PR 30-MAR-2001; 2001US-0279766P.
XX
PA (FARB) BAYER AG.
XX
PI Xiao Y;
XX
DR WPI; 2002-122283/16.
XX
PT Novel purified human eosinophil serine protease 1-like enzyme, useful for
PT identifying modulators of enzyme activity for treating Paget's disease,
PT osteoporosis, airway allergy, asthma.
XX
PS Disclosure; Fig 4; 131pp; English.
XX
CC The invention relates to a purified human eosinophil serine protease-1
CC (esp-1) like enzyme. Esp-1 like enzyme is useful in in-vitro or in-vivo

```
CC   assays to identify test compounds with potential therapeutic or
CC   diagnostic value. Esp1-like enzyme modulator is useful for treating esp1-
CC   like enzyme dysfunction related diseases condition such as asthma, airway
CC   allergy, chronic obstructive pulmonary disease (COPD) or osteoporosis.
CC   Esp-1 like enzyme is also useful in diagnostic assays for detecting
CC   diseases and abnormalities or susceptibility to diseases related to
CC   presence of mutations in the nucleic acid sequences which encode the
CC   enzyme. Pharmaceutical composition comprising esp-1 like enzyme is useful
CC   for treating dermatitis, Paget's disease, and preventing degradation of
CC   bone implants particularly dental implants. The present sequence is human
CC   esp-1 like enzyme
XX
SQ   Sequence 314 AA;

  Query Match             99.7%;  Score 1723;  DB 5;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1.7e-146;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314


RESULT 11
ABU71634
ID   ABU71634 standard; protein; 314 AA.
XX
AC   ABU71634;
XX
DT   16-JUN-2003  (first entry)
XX
DE   Human PRO polypeptide #45.
XX
KW   Human; PRO; secreted polypeptide; transmembrane polypeptide;
KW   pathological disorder; cardiac insufficiency disorder; protein secretion;
KW   pancreas; diabetes; gastrointestinal mucosa; mucosal lesion; psoriasis;
KW   skin disease; keratinocyte differentiation; epithelial cancer; tumour;
KW   lung squamous cell carcinoma; epidermoid carcinoma; vulva; glioma;
KW   cytostatic; cardiant; endocrine; antidiabetic; gastrointestinal;
KW   antiulcer; dermatological; vulnerary.
XX
OS   Homo sapiens.
XX
PN   US2002146709-A1.
XX
PD   10-OCT-2002.
XX
PF   18-JUL-2001; 2001US-00909088.
XX
PR   17-SEP-1997;   97US-0059113P.
PR   17-SEP-1997;   97US-0059115P.
PR   17-SEP-1997;   97US-0059117P.
PR   17-SEP-1997;   97US-0059119P.
PR   17-SEP-1997;   97US-0059121P.
PR   17-SEP-1997;   97US-0059122P.
PR   17-SEP-1997;   97US-0059184P.
PR   18-SEP-1997;   97US-0059263P.
PR   18-SEP-1997;   97US-0059266P.
PR   15-OCT-1997;   97US-0062125P.
PR   17-OCT-1997;   97US-0062285P.
PR   17-OCT-1997;   97US-0062287P.
PR   21-OCT-1997;   97US-0063486P.
PR   24-OCT-1997;   97US-0062814P.
PR   24-OCT-1997;   97US-0062816P.
PR   24-OCT-1997;   97US-0063045P.
PR   24-OCT-1997;   97US-0063120P.
PR   24-OCT-1997;   97US-0063121P.
PR   24-OCT-1997;   97US-0063127P.
PR   24-OCT-1997;   97US-0063128P.
PR   27-OCT-1997;   97US-0063327P.
PR   27-OCT-1997;   97US-0063329P.
PR   28-OCT-1997;   97US-0063541P.
PR   28-OCT-1997;   97US-0063542P.
PR   28-OCT-1997;   97US-0063544P.
PR   28-OCT-1997;   97US-0063549P.
PR   28-OCT-1997;   97US-0063550P.
PR   28-OCT-1997;   97US-0063564P.
PR   29-OCT-1997;   97US-0063435P.
PR   29-OCT-1997;   97US-0063704P.
PR   29-OCT-1997;   97US-0063732P.
PR   29-OCT-1997;   97US-0063734P.
PR   29-OCT-1997;   97US-0063735P.
PR   29-OCT-1997;   97US-0063738P.
PR   29-OCT-1997;   97US-0064215P.
PR   31-OCT-1997;   97US-0063870P.
PR   31-OCT-1997;   97US-0064103P.
PR   03-NOV-1997;   97US-0064248P.
PR   07-NOV-1997;   97US-0064809P.
PR   12-NOV-1997;   97US-0065186P.
PR   17-NOV-1997;   97US-0065846P.
PR   18-NOV-1997;   97US-0065693P.
PR   21-NOV-1997;   97US-0066120P.
PR   21-NOV-1997;   97US-0066364P.
PR   24-NOV-1997;   97US-0066453P.
PR   24-NOV-1997;   97US-0066466P.
PR   24-NOV-1997;   97US-0066511P.
PR   24-NOV-1997;   97US-0066770P.
PR   24-NOV-1997;   97US-0066772P.
PR   10-SEP-1998;   98WO-US018824.
PR   14-SEP-1998;   98WO-US019177.
PR   16-SEP-1998;   98WO-US019330.
PR   17-SEP-1998;   98WO-US019437.
PR   01-DEC-1998;   98WO-US025108.
PR   08-SEP-1999;   99WO-US020594.
PR   13-SEP-1999;   99WO-US020944.
PR   15-SEP-1999;   99WO-US021090.
PR   15-SEP-1999;   99WO-US021547.
PR   05-OCT-1999;   99WO-US023089.
PR   29-NOV-1999;   99WO-US028214.
PR   30-NOV-1999;   99WO-US028313.
PR   01-DEC-1999;   99WO-US028301.
PR   02-DEC-1999;   99WO-US028564.
PR   02-DEC-1999;   99WO-US028565.
PR   16-DEC-1999;   99WO-US030095.
PR   20-DEC-1999;   99WO-US030911.
PR   20-DEC-1999;   99WO-US030999.
PR   05-JAN-2000; 2000WO-US000219.
PR   11-FEB-2000; 2000WO-US003565.
PR   22-FEB-2000; 2000WO-US004414.
PR   24-FEB-2000; 2000WO-US005004.
PR   02-MAR-2000; 2000WO-US005841.
PR   20-MAR-2000; 2000WO-US007377.
PR   30-MAR-2000; 2000WO-US008439.
PR   22-MAY-2000; 2000WO-US014042.
PR   02-JUN-2000; 2000WO-US015264.
PR   28-JUL-2000; 2000WO-US020710.
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PR   24-AUG-2000; 2000WO-US023328.
PR   18-SEP-2000; 2000US-00665350.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Ashkenazi A,  Botstein D,  Desnoyers L,  Eaton DL,  Ferrara N;
PI   Filvaroff E,  Fong S,  Gao W,  Gerber H,  Gerritsen ME,  Goddard A;
PI   Godowski PJ,  Grimaldi JC,  Gurney AL,  Hillan KJ,  Kljavin IJ;
PI   Mather JP,  Pan J,  Paoni NF,  Roy MA,  Stewart TA,  Tumas D;
PI   Williams PM,  Wood WI;
XX
DR   WPI; 2003-328338/31.
DR   N-PSDB; ACA59107.
XX
PT   Isolated nucleic acid useful for e.g., treating pathological disorders
PT   encodes a secreted or transmembrane protein.
XX
PS   Claim 12; Fig 92; 473pp; English.
XX
CC   The invention relates to human PRO polypeptides (secreted or
CC   transmembrane polypeptides) and the polynucleotides encoding them. The
CC   PRO polypeptides and polynucleotides can be used in treating pathological
CC   disorders and tumours, in therapeutic treatment of cardiac insufficiency
CC   disorders and in therapeutic treatment of disorders involving protein
CC   secretion by the pancreas, including diabetes. They can also be used in
CC   treating disorders associated with the preservation and maintenance of
CC   gastrointestinal mucosa and the repair of acute and chronic mucosal
CC   lesions, and skin diseases associated with abnormal keratinocyte
CC   differentiation (e.g., psoriases, epithelial cancers such as lung
CC   squamous cell carcinoma, epidermoid carcinoma of the vulva and gliomas).
CC   The sequences can be used as molecular markers for protein
CC   electrophoresis purposes and can be utilised in protein-protein binding
CC   assays, biochemical screening assays, immunoassays and cell-based assays.
CC   This sequence represents a human PRO polypeptide of the invention
XX
SQ   Sequence 314 AA;

  Query Match             99.7%;  Score 1723;  DB 6;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1.7e-146;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314


RESULT 12
ABU71489
ID   ABU71489 standard; protein; 314 AA.
XX
AC   ABU71489;
XX
DT   10-JUN-2003  (first entry)
XX
DE   Human PRO polypeptide #45.
XX
KW   Human; secreted and transmembrane protein; PRO polypeptide; cancer;
KW   Alzheimer's disease; ischaemia; cytostatic; nootropic; vasotropic;
KW   neuroprotective.
XX
OS   Homo sapiens.
XX
PN   US2002192659-A1.
XX
PD   19-DEC-2002.
XX
PF   10-JUL-2001; 2001US-00902853.
XX
PR   17-SEP-1997;   97US-0059113P.
PR   17-SEP-1997;   97US-0059115P.
PR   17-SEP-1997;   97US-0059117P.
PR   17-SEP-1997;   97US-0059119P.
PR   17-SEP-1997;   97US-0059121P.
PR   17-SEP-1997;   97US-0059122P.
PR   17-SEP-1997;   97US-0059184P.
PR   18-SEP-1997;   97US-0059263P.
PR   18-SEP-1997;   97US-0059266P.
PR   15-OCT-1997;   97US-0062125P.
PR   17-OCT-1997;   97US-0062285P.
PR   17-OCT-1997;   97US-0062287P.
PR   21-OCT-1997;   97US-0063486P.
PR   24-OCT-1997;   97US-0062814P.
PR   24-OCT-1997;   97US-0062816P.
PR   24-OCT-1997;   97US-0063045P.
PR   24-OCT-1997;   97US-0063120P.
PR   24-OCT-1997;   97US-0063121P.
PR   24-OCT-1997;   97US-0063127P.
PR   24-OCT-1997;   97US-0063128P.
PR   27-OCT-1997;   97US-0063327P.
PR   27-OCT-1997;   97US-0063329P.
PR   28-OCT-1997;   97US-0063541P.
PR   28-OCT-1997;   97US-0063542P.
PR   28-OCT-1997;   97US-0063544P.
PR   28-OCT-1997;   97US-0063549P.
PR   28-OCT-1997;   97US-0063550P.
PR   28-OCT-1997;   97US-0063564P.
PR   29-OCT-1997;   97US-0063435P.
PR   29-OCT-1997;   97US-0063704P.
PR   29-OCT-1997;   97US-0063732P.
PR   29-OCT-1997;   97US-0063734P.
PR   29-OCT-1997;   97US-0063735P.
PR   29-OCT-1997;   97US-0063738P.
PR   29-OCT-1997;   97US-0064215P.
PR   31-OCT-1997;   97US-0063870P.
PR   31-OCT-1997;   97US-0064103P.
PR   03-NOV-1997;   97US-0064248P.
PR   07-NOV-1997;   97US-0064809P.
PR   12-NOV-1997;   97US-0065186P.
PR   17-NOV-1997;   97US-0065846P.
PR   18-NOV-1997;   97US-0065693P.
PR   21-NOV-1997;   97US-0066120P.
PR   21-NOV-1997;   97US-0066364P.
PR   24-NOV-1997;   97US-0066453P.
PR   24-NOV-1997;   97US-0066466P.
PR   24-NOV-1997;   97US-0066511P.
PR   24-NOV-1997;   97US-0066770P.
PR   24-NOV-1997;   97US-0066772P.
PR   10-SEP-1998;   98WO-US018824.
PR   14-SEP-1998;   98WO-US019177.
PR   16-SEP-1998;   98WO-US019330.
PR   17-SEP-1998;   98WO-US019437.
PR   01-DEC-1998;   98WO-US025108.
PR   08-SEP-1999;   99WO-US020594.
PR   13-SEP-1999;   99WO-US020944.
PR   15-SEP-1999;   99WO-US021090.
PR   15-SEP-1999;   99WO-US021547.
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PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-361832/34.
DR N-PSDB; ACA58504.
XX
PT New isolated nucleic acid encoding a PRO polypeptide, e.g. PRO245 or
PT PRO1868, useful in molecular biology, chromosome and gene mapping, in
PT generating antisense RNA and DNA, and in gene therapy.
XX
PS Claim 12; Fig 92; 474pp; English.
XX
CC The present invention relates to the isolation of novel human secreted
CC and transmembrane proteins (PRO polypeptides), and the polynucleotide
CC sequences encoding them. The polynucleotide sequences are useful in
CC molecular biology, as hybridisation probes, in chromosome and gene
CC mapping, in generating antisense RNA and DNA, and in gene therapy. The
CC polynucleotide sequences may also be used in preparing PRO polypeptides
CC by recombinant techniques, and in generating either transgenic animals or
CC knock-out animals which, in turn, are useful in the development and
CC screening of therapeutically useful reagents. The PRO polypeptides or
CC their antibodies are useful in preparing a medicament for treating a
CC condition responsive to the polypeptide or antibody, such as cancer,
CC Alzheimer's disease or ischaemia, and in various diagnostic assays.
CC ABU71445-ABU71505 represent human PRO polypeptides of the invention
XX
SQ Sequence 314 AA;

Query Match 99.7%; Score 1723; DB 6; Length 314;
Best Local Similarity 99.7%; Pred. No. 1.7e-146;
Matches 313; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

```
Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314
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RESULT 13
ABU71935
ID ABU71935 standard; protein; 314 AA.
XX
AC ABU71935;
XX
DT 12-JUN-2003 (first entry)
XX
DE Human secreted/transmembrane protein PRO303.
XX
KW Human; secreted protein; transmembrane protein; PRO; gene therapy;
KW chromosome identification; chromosome marker.
XX
OS Homo sapiens.
XX
PN US2003003530-A1.
XX
PD 02-JAN-2003.
XX
PF 11-JUL-2001; 2001US-00904011.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
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PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.

PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 01-DEC-1998; 98WO-US025108.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-329602/31.
DR N-PSDB; ACA60211.
XX
PT New transmembrane polypeptides and nucleic acids encoding the
PT polypeptides, useful in gene therapy, in chromosome identification, as
PT chromosome markers, in generating probes and in tissue typing.
XX
PS Claim 12; Fig 92; 484pp; English.
XX
CC The invention relates to an isolated nucleic acid with at least 80%
CC nucleic acid sequence identity to a nucleotide sequence encoding one of
CC 61 secreted/transmembrane polypeptides, or PRO polypeptides or encoding a
CC PRO protein extracellular domain. Also included are a vector comprising
CC the PRO nucleic acid, a host cell comprising the vector, producing a PRO
CC polypeptide (by culturing the host cell for the expression of the PRO
CC polypeptide, and recovering the PRO polypeptide from the cell culture),
CC an isolated PRO polypeptide (having at least 80% sequence identity to: (
CC a) an amino acid sequence selected from the 61 PRO proteins; (b) an amino
CC acid sequence encoded by a nucleic acid molecule deposited with an ATCC
CC number (detailed in the specification); or (c) an extracellular domain of
CC a PRO polypeptide or to a PRO polypeptide lacking its associated signal
CC peptide), a chimaeric molecule comprising a PRO polypeptide of fused to a
CC heterologous amino acid sequence, an anti-PRO antibody, detecting a
CC PRO245 or PRO1868 in a sample suspected of containing the polypeptide,
CC linking a bioactive molecule to a cell expressing a PRO245 or PRO1868 and
CC modulating at least one biological activity of a cell expressing a PRO245
CC or PRO1868. Nucleic acids which encode PRO can be used to generate either
CC transgenic animals or knock-out animals which may be used in the
CC development and screening of therapeutically useful reagents. The nucleic
CC acids may also be used in gene therapy, in chromosome identification, as
CC chromosome markers, or in generating probes. The PRO polypeptides are
CC useful as molecular markers for protein electrophoresis, and the isolated
CC nucleic acids may be used for recombinantly expressing those markers. The
CC PRO polypeptides and nucleic acids may also be used in tissue typing.
CC Anti-PRO antibodies are useful in diagnostic assays for PRO, and in
CC affinity purification of PRO from recombinant cell culture or natural
CC sources. The present sequence represents a PRO protein
XX
SQ Sequence 314 AA;

Query Match 99.7%; Score 1723; DB 6; Length 314;
Best Local Similarity 99.7%; Pred. No. 1.7e-146;
Matches 313; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

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Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314
```

RESULT 14
ABO01818
ID ABO01818 standard; protein; 314 AA.
XX
AC ABO01818;
XX
DT 07-AUG-2003 (first entry)
XX
DE Novel human secreted and transmembrane protein PRO303.
XX
KW Human; secreted and transmembrane protein; PRO; pharmaceutical;
KW diagnostic; biosensor; bioreactor; Parkinson's disease;
KW Alzheimer's disease; inflammation; nephritis; wound healing;
KW nerve repair; collateral blood vessel formation; cancer;
KW colorectal cancer; haemorrhage; rheumatoid arthritis; diabetes;
KW cirrhosis; fibrosis; restenosis; dermal fibrotic condition; keloid;
KW scarring; ischaemia; stroke; hypertension; heart attack; atherosclerosis;
KW infertility; gene therapy.
XX
OS Homo sapiens.
XX
PN US2002197671-A1.
XX
PD 26-DEC-2002.
XX
PF 17-JUL-2001; 2001US-00907824.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.

PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 01-DEC-1998; 98WO-US025108.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-370793/35.
DR N-PSDB; ACD07611.
XX
PT New genes and secreted and transmembrane polypeptides (e.g. PRO245 or
PT PRO335), useful for treating or diagnosing e.g. Alzheimer's disease,
PT cancers, hemorrhage, rheumatoid arthritis, diabetes, cirrhosis, ischemia
PT or strokes.
XX
PS Claim 12; Fig 92; 482pp; English.
XX
CC The invention describes a new isolated nucleic acid molecule comprising
CC the full length coding sequence of the DNA deposited with the American
CC Type Culture Collection (e.g. ATCC Deposit No. 209258) ,or a sequence
CC with at least 80% identity to a DNA encoding a PRO polypeptide comprising
CC any of 61 sequences having 164-1119 amino acids fully defined in the
CC specification. The PRO polypeptides or polynucleotides are useful as
CC pharmaceuticals, diagnostics, biosensors or bioreactors. These are
CC particularly useful for detecting or treating e.g. Parkinson's disease,
CC Alzheimer's disease, inflammations, nephritis, wound healing, nerve
CC repair, collateral blood vessel formation, cancers (e.g. colorectal
CC cancer), haemorrhage (or reduce risk for haemorrhage), rheumatoid
CC arthritis, diabetes, cirrhosis of the liver, fibrosis of the lungs,
CC restenosis, dermal fibrotic conditions (e.g. keloids or scarring),
CC ischaemia, strokes, hypertension, heart attacks, atherosclerosis, or
CC infertility in mammals (e.g. humans, dogs, cats, cattle, horses, sheep,
CC pigs, goats, or rabbits) The PRO polypeptides are useful as targets for
CC therapeutic intervention in these diseases, and diagnostic determination
CC of the presence of these diseases. The PRO polypeptides are also useful
CC as molecular weight markers, or for chromosome identification. The PRO
CC genes are useful as hybridisation probes, or for screening libraries of
CC human cDNA, genomic DNA or mRNA. The PRO genes may also be used in gene
CC therapy, particularly for replacing a defective gene. This is the amino
CC acid sequence of a novel human secreted and transmembrane PRO polypeptide
XX
SQ Sequence 314 AA;

Query Match 99.7%; Score 1723; DB 6; Length 314;
Best Local Similarity 99.7%; Pred. No. 1.7e-146;
Matches 313; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

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Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
```

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                  ||||||||||||||||
Db        301 PFPLLWALPLLGPV 314

RESULT 15
ABU54391
ID   ABU54391 standard; protein; 314 AA.
XX
AC   ABU54391;
XX
DT   10-MAR-2003  (first entry)
XX
DE   Human secreted/transmembrane protein PRO303.
XX
KW   Human; PRO; secreted protein; transmembrane protein; enterocolitis;
KW   gastrointestinal ulceration; skin disease;
KW   abnormal keratinocyte differentiation; psoriasis; epithelial cancer;
KW   squamous cell carcinoma; Alzheimer's disease; Parkinson's disease;
KW   amyotrophic lateral sclerosis; inflammatory disease;
KW   rheumatoid arthritis; asthma; multiple sclerosis; organ failure;
KW   atherosclerosis; cardiac injury; infertility; birth defect;
KW   premature aging; AIDS; acquired immunodeficiency syndrome; cancer;
KW   diabetic complication; wound repair.
XX
OS   Homo sapiens.
XX
PN   US2002132240-A1.
XX
PD   19-SEP-2002.
XX
PF   18-JUL-2001; 2001US-00909320.
XX
PR   17-SEP-1997;   97US-0059113P.
PR   17-SEP-1997;   97US-0059115P.
PR   17-SEP-1997;   97US-0059117P.
PR   17-SEP-1997;   97US-0059119P.
PR   17-SEP-1997;   97US-0059121P.
PR   17-SEP-1997;   97US-0059122P.
PR   17-SEP-1997;   97US-0059184P.
PR   18-SEP-1997;   97US-0059263P.
PR   18-SEP-1997;   97US-0059266P.
PR   15-OCT-1997;   97US-0062125P.
PR   17-OCT-1997;   97US-0062285P.
PR   17-OCT-1997;   97US-0062287P.
PR   21-OCT-1997;   97US-0063486P.
PR   24-OCT-1997;   97US-0062814P.
PR   24-OCT-1997;   97US-0062816P.
PR   24-OCT-1997;   97US-0063045P.
PR   24-OCT-1997;   97US-0063120P.
PR   24-OCT-1997;   97US-0063121P.
PR   24-OCT-1997;   97US-0063127P.
PR   24-OCT-1997;   97US-0063128P.
PR   27-OCT-1997;   97US-0063327P.
PR   27-OCT-1997;   97US-0063329P.
PR   28-OCT-1997;   97US-0063541P.
PR   28-OCT-1997;   97US-0063542P.
PR   28-OCT-1997;   97US-0063544P.
PR   28-OCT-1997;   97US-0063549P.
PR   28-OCT-1997;   97US-0063550P.
PR   28-OCT-1997;   97US-0063564P.
PR   29-OCT-1997;   97US-0063435P.
PR   29-OCT-1997;   97US-0063704P.
PR   29-OCT-1997;   97US-0063732P.
PR   29-OCT-1997;   97US-0063734P.
PR   29-OCT-1997;   97US-0063735P.
PR   29-OCT-1997;   97US-0063738P.
PR   29-OCT-1997;   97US-0064215P.
PR   31-OCT-1997;   97US-0063870P.
PR   31-OCT-1997;   97US-0064103P.
PR   03-NOV-1997;   97US-0064248P.
PR   07-NOV-1997;   97US-0064809P.
PR   12-NOV-1997;   97US-0065186P.
PR   17-NOV-1997;   97US-0065846P.
PR   18-NOV-1997;   97US-0065693P.
PR   21-NOV-1997;   97US-0066120P.
PR   21-NOV-1997;   97US-0066364P.
PR   24-NOV-1997;   97US-0066453P.
PR   24-NOV-1997;   97US-0066466P.
PR   24-NOV-1997;   97US-0066511P.
PR   24-NOV-1997;   97US-0066770P.
PR   24-NOV-1997;   97US-0066772P.
PR   10-SEP-1998;   98WO-US018824.
PR   14-SEP-1998;   98WO-US019177.
PR   16-SEP-1998;   98WO-US019330.
PR   17-SEP-1998;   98WO-US019437.
PR   01-DEC-1998;   98WO-US025108.
PR   08-SEP-1999;   99WO-US020594.
PR   13-SEP-1999;   99WO-US020944.
PR   15-SEP-1999;   99WO-US021090.
PR   15-SEP-1999;   99WO-US021547.
PR   05-OCT-1999;   99WO-US023089.
PR   29-NOV-1999;   99WO-US028214.
PR   30-NOV-1999;   99WO-US028313.
PR   01-DEC-1999;   99WO-US028301.
PR   02-DEC-1999;   99WO-US028564.
PR   02-DEC-1999;   99WO-US028565.
PR   16-DEC-1999;   99WO-US030095.
PR   20-DEC-1999;   99WO-US030911.
PR   20-DEC-1999;   99WO-US030999.
PR   06-JAN-2000; 2000WO-US000219.
PR   11-FEB-2000; 2000WO-US003565.
PR   22-FEB-2000; 2000WO-US004414.
PR   24-FEB-2000; 2000WO-US005004.
PR   02-MAR-2000; 2000WO-US005841.
PR   20-MAR-2000; 2000WO-US007377.
PR   30-MAR-2000; 2000WO-US008439.
PR   22-MAY-2000; 2000WO-US014042.
PR   02-JUN-2000; 2000WO-US015264.
PR   28-JUL-2000; 2000WO-US020710.
PR   24-AUG-2000; 2000WO-US023328.
PR   18-SEP-2000; 2000US-00665350.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Ashkenazi A,  Botstein D,  Desnoyers L,  Eaton DL,  Ferrara N;
PI   Filvaroff E,  Fong S,  Gao W,  Gerber H,  Gerritsen ME,  Goddard A;
PI   Godowski PJ,  Grimaldi JC,  Gurney AL,  Hillan KJ,  Kljavin IJ;
PI   Mather JP,  Pan J,  Paoni NF,  Roy MA,  Stewart TA,  Tumas D;
PI   Williams PM,  Wood WI;
XX
DR   WPI; 2003-147434/14.
DR   N-PSDB; ABX71659.
XX
PT   New PRO polypeptides and nucleic acid molecules, useful in diagnosing or
PT   treating inflammatory diseases, organ failure, atherosclerosis, cardiac
PT   injury, infertility, cancer, AIDS, Alzheimer's disease or Parkinson's
PT   disease.
XX
PS   Claim 12; Fig 92; 473pp; English.
XX
CC   The invention relates to an isolated PRO polypeptide having at least 80%
CC   amino acid sequence identity to: (a) any one of 61 fully defined amino
CC   acid sequences given in the specification (appearing as ABU54347-
CC   ABU54407); (b) an amino acid sequence encoded by the nucleotide sequence
CC   deposited under American Type Culture Collection (accession numbers
CC   listed in the specification); (c) any one of the PRO sequences which
CC   lacks its associated signal peptide; (d) an extracellular domain of the
CC   PRO polypeptide with its associated signal peptide; or (e) an
CC   extracellular domain of the PRO polypeptide which lacks its associated
CC   signal peptide. Also include are the nucleic acids encoding the PRO
CC   polypeptides, vectors, host cells and anti-PRO antibodies. The PRO
CC   polypeptides and nucleic acids are useful in diagnosing or treating
CC   enterocolitis, gastrointestinal ulceration, skin diseases associated with
CC   abnormal keratinocyte differentiation, e.g. psoriasis or epithelial
CC   cancers such as squamous cell carcinoma, Alzheimer's disease, Parkinson's
CC   disease, amyotrophic lateral sclerosis, inflammatory diseases, e.g.
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CC   rheumatoid arthritis, asthma or multiple sclerosis, organ failure,
CC   atherosclerosis, cardiac injury, infertility, birth defects, premature
CC   aging, AIDS, cancer, diabetic complications, or mutations in general. The
CC   polypeptides are also useful for wound repair and associated therapies
CC   concerned with re-growth of tissue. The nucleotide sequences may be used
CC   as hybridisation probes in chromosome and gene mapping, or in generating
CC   antisense RNA and DNA. PRO nucleic acids are also useful in preparing PRO
CC   polypeptides, in assays to identify other proteins or molecules involved
CC   in binding reaction, to generate transgenic animals or knockout animals,
CC   which in turn are useful in the development and screening of
CC   therapeutically useful reagents, for chromosome identification, and
CC   tissue typing. The PRO polypeptides and nucleic acid molecules are also
CC   useful in gene therapy, and as molecular weight markers for protein
CC   electrophoresis purposes. The anti-PRO antibodies may be used in
CC   diagnostic assays for PRO, or for the affinity purification of PRO from
CC   recombinant cell culture or natural sources. The present sequence
CC   represents a PRO polypeptide
XX
SQ   Sequence 314 AA;

  Query Match             99.7%;  Score 1723;  DB 6;  Length 314;
  Best Local Similarity   99.7%;  Pred. No. 1.7e-146;
  Matches  313;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MGARGALLLALLLARAGLRKPESQEAAPLSGPCGRRVITSRIVGGEDAELGRWPWQGSLR 60

Qy         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 LWDSHVCGVSLLSHRWALTAAHCFETYSDLSDPSGWMVQFGQLTSMPSFWSLQAYYTRYF 120

Qy        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 VSNIYLSPRYLGNSPYDIALVKLSAPVTYTKHIQPICLQASTFEFENRTDCWVTGWGYIK 180

Qy        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 EDEALPSPHTLQEVQVAIINNSMCNHLFLKYSFRKDIFGDMVCAGNAQGGKDACFGDSGG 240

Qy        241 PLACNKDGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300
              ||||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 PLACNKNGLWYQIGVVSWGVGCGRPNRPGVYTNISHHFEWIQKLMAQSGMSQPDPSWPLL 300

Qy        301 FFPLLWALPLLGPV 314
              ||||||||||||||
Db        301 FFPLLWALPLLGPV 314


Search completed: February 27, 2005, 20:22:32
Job time : 118.374 secs
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GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: March 4, 2005, 22:14:12 ; Search time 4246.77 Seconds
(without alignments)
9859.425 Million cell updates/sec

Title: US-10-040-647-5
Perfect score: 1100
Sequence: 1 CGCGGGAGAGGAGGCCATGG.....AAAAAAAAAAAAAAAAAAAA 1100

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 34239544 seqs, 19032134700 residues

Total number of hits satisfying chosen parameters: 68479088

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : EST:*
1: gb_est1:*
2: gb_est2:*
3: gb_htc:*
4: gb_est3:*
5: gb_est4:*
6: gb_est5:*
7: gb_est6:*
8: gb_gss1:*
9: gb_gss2:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Result No.	Score	% Query Match	Length	DB	ID	Description
	1	1071.8	97.4	1077	3	CR590593	CR590593 full-leng
	2	1012.4	92.0	1014	3	CR605277	CR605277 full-leng
c	3	908.4	82.6	932	1	AL578261	AL578261
	4	870.4	79.1	884	1	AL555870	AL555870
	5	866	78.7	1057	5	BX417595	BX417595
c	6	780.6	71.0	898	5	BX454526	BX454526
	7	777.2	70.7	871	4	BI334059	BI334059 602997320
	8	760.4	69.1	777	4	BI259237	BI259237 602970095
	9	733.6	66.7	878	4	BI827608	BI827608 603073575
c	10	669.8	60.9	743	6	CB529999	CB529999 UI-H-FT2-
c	11	624.8	56.8	633	1	AI126185	AI126185 qd81h05.x
	12	569.2	51.7	705	4	BM019026	BM019026 603647010
	13	550.4	50.0	682	4	BM017444	BM017444 603644544
	14	548.8	49.9	1219	4	BM553333	BM553333 AGENCOURT
c	15	525.8	47.8	591	1	AA161187	AA161187 zo59b06.s
	16	524.8	47.7	824	4	BM018961	BM018961 603646918
	17	519	47.2	1286	5	BM911057	BM911057 AGENCOURT
	18	500.6	45.5	825	4	BI257649	BI257649 602967571
	19	489.4	44.5	1050	3	AK006271	AK006271 Mus muscu
	20	482.6	43.9	624	4	BM018436	BM018436 603646140
	21	476.4	43.3	625	4	BG717268	BG717268 602689582
c	22	473.6	43.1	514	1	AA620757	AA620757 af90d04.s
	23	472.4	42.9	628	4	BG717347	BG717347 602689681
	24	464.8	42.3	479	1	AA448095	AA448095 zw82c04.r
	25	444.2	40.4	510	4	BM759973	BM759973 K-EST0040
c	26	441.6	40.1	462	1	AI016399	AI016399 ov59h09.s
	27	440.4	40.0	463	7	CR735731	CR735731 CR735731
c	28	418.4	38.0	487	1	AA525010	AA525010 nh36a02.s
	29	414.2	37.7	1018	6	BY706352	BY706352 BY706352
c	30	401.2	36.5	406	1	AI150137	AI150137 qf44g06.x
c	31	396.2	36.0	412	1	AA448166	AA448166 zw83a10.s
c	32	395.6	36.0	402	1	AA729680	AA729680 nx37d02.s
c	33	394.2	35.8	399	1	AI278269	AI278269 ql78b05.x
c	34	371.6	33.8	383	1	AA447596	AA447596 zw82c04.s
c	35	362.4	32.9	670	6	CB530049	CB530049 UI-H-FT2-
c	36	362.4	32.9	677	6	CD365369	CD365369 UI-H-FT2-
	37	352.8	32.1	454	4	BI257498	BI257498 602967618
c	38	349.4	31.8	373	1	AA401115	AA401115 zu50g12.s
	39	337	30.6	1089	3	AK016509	AK016509 Mus muscu
c	40	336.8	30.6	475	2	AW971363	AW971363 EST383452
c	41	329	29.9	411	1	AA149942	AA149942 zo68d04.s
	42	324	29.5	545	6	CB158100	CB158100 K-EST0217
c	43	316.6	28.8	334	1	AA552523	AA552523 nk14e05.s
c	44	309.6	28.1	351	1	AA860356	AA860356 aj59b05.s
	45	308.8	28.1	766	6	CA465071	CA465071 AGENCOURT

ALIGNMENTS

RESULT 1
CR590593
LOCUS CR590593 1077 bp mRNA linear HTC 21-JUL-2004
DEFINITION full-length cDNA clone CS0DE010YP05 of Placenta of Homo sapiens (human).
ACCESSION CR590593
VERSION CR590593.1 GI:50471400
KEYWORDS HTC; CNSLT_cDNA.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 1077)
AUTHORS Li,W.B., Gruber,C., Jessee,J. and Polayes,D.
TITLE Full-length cDNA libraries and normalization
JOURNAL Unpublished
REMARK Contact : Feng Liang Email : fliang@lifetech.com URL : http://fulllength.invitrogen.com/ InVitroGen Corporation 1600 Faraday Avenue
REFERENCE 2 (bases 1 to 1077)
AUTHORS Genoscope.
TITLE Direct Submission
JOURNAL Submitted (20-JUL-2004) Genoscope - Centre National de Sequencage : BP 191 91006 EVRY cedex - FRANCE (E-mail : seqref@genoscope.cns.fr - Web : www.genoscope.cns.fr)
COMMENT 1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-strand cDNA was digested with Not I and cloned into the Not I and EcoR V sites of the pCMVSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies, a division of Invitrogen.
FEATURES Location/Qualifiers
source 1. .1077
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="CS0DE010YP05"
/tissue_type="Placenta"
/plasmid="pCMVSPORT_6"
ORIGIN

Query Match 97.4%; Score 1071.8; DB 3; Length 1077;
Best Local Similarity 99.8%; Pred. No. 1.5e-258;
Matches 1073; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 3 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 62

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Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         63 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 122

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        123 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 182

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        183 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 242

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        243 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 302

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        303 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 362

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        363 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 422

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        423 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 482

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        483 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 542

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        543 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 602

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Db        603 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 662

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        663 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 722

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        723 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 782

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        783 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 842

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        843 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 902

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        903 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 962

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        963 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1022

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAA 1075
              ||||||||||||||||||||||||||||||||||||||||||||| |||||||
Db       1023 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAA 1077
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RESULT 2
CR605277
LOCUS CR605277 1014 bp mRNA linear HTC 21-JUL-2004

DEFINITION full-length cDNA clone CS0DK002YM07 of HeLa cells Cot 25-normalized of Homo sapiens (human).
ACCESSION CR605277
VERSION CR605277.1 GI:50486084
KEYWORDS HTC; CNSLT_cDNA.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 1014)
AUTHORS Li,W.B., Gruber,C., Jessee,J. and Polayes,D.
TITLE Full-length cDNA libraries and normalization
JOURNAL Unpublished
REMARK Contact : Feng Liang Email : fliang@lifetech.com URL : http://fulllength.invitrogen.com/ InVitroGen Corporation 1600 Faraday Avenue
REFERENCE 2 (bases 1 to 1014)
AUTHORS Genoscope.
TITLE Direct Submission
JOURNAL Submitted (20-JUL-2004) Genoscope - Centre National de Sequencage : BP 191 91006 EVRY cedex - FRANCE (E-mail : seqref@genoscope.cns.fr - Web : www.genoscope.cns.fr)
COMMENT 1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-strand cDNA was digested with Not I and cloned into the Not I and EcoR V sites of the pCMVSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies, a division of Invitrogen.
FEATURES Location/Qualifiers
source 1. .1014
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="CS0DK002YM07"
/tissue_type="HeLa cells Cot 25-normalized"
/plasmid="pCMVSPORT_6"
ORIGIN

Query Match 92.0%; Score 1012.4; DB 3; Length 1014;
Best Local Similarity 99.9%; Pred. No. 1.3e-243;
Matches 1013; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy          3 CGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCGGG 62
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 CGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCGGG 60

Qy         63 CTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCGAC 122
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 CTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCGAC 120

Qy        123 GGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTGGC 182
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 GGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTGGC 180

Qy        183 AGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGCT 242
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 AGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGCT 240

Qy        243 GGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGGGT 302
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 GGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGGGT 300

Qy        303 GGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTACT 362
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Db        301 GGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTACT 360

Qy        363 ACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACCCT 422
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Db        361 ACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACCCT 420

Qy        423 ATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCCCA 482
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Db        421 ATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCCCA 480
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Qy        483 TCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCT 542
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Db        481 TCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCT 540

Qy        543 GGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCAGG 602
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Db        541 GGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCAGG 600

Qy        603 TCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGG 662
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Db        601 TCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGG 660

Qy        663 ACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTCG 722
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        661 ACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTCG 720

Qy        723 GTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGTCG 782
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Db        721 GTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGTCG 780

Qy        783 TGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGCC 842
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Db        781 TGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGCC 840

Qy        843 ACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCCCT 902
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Db        841 ACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCCCT 900

Qy        903 CCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTGAG 962
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        901 CCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTGAG 960

Qy        963 CCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCT 1016
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Db        961 CCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCT 1014
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RESULT 3
AL578261/c
LOCUS AL578261 932 bp mRNA linear EST 07-APR-2004
DEFINITION AL578261 Homo sapiens HELA CELLS COT 25-NORMALIZED Homo sapiens cDNA clone CS0DK002YM07 3-PRIME, mRNA sequence.
ACCESSION AL578261
VERSION AL578261.3 GI:46257165
KEYWORDS EST.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 932)
AUTHORS Li,W.B., Gruber,C., Jessee,J. and Polayes,D.
TITLE Full-length cDNA libraries and normalization
JOURNAL Unpublished (2001)
COMMENT On Feb 16, 2001 this sequence version replaced gi:31316478.
Contact: Genoscope
Genoscope - Centre National de Sequencage
2 rue Gaston Cremieux, CP 5706 - 91057 EVRY cedex - FRANCE
Email: seqref@genoscope.cns.fr, Web : www.genoscope.cns.fr
1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-strand cDNA was digested with Not I and cloned into the Not I and EcoR V sites of the pCMVSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies, a division of Invitrogen. This sequence belongs to sequence cluster 8290.f
For more information about this cluster, see
http://www.genoscope.cns.fr/cdna?s=CS0DK002AG04NP1&c=8290.f.
FEATURES Location/Qualifiers
source 1. .932
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="CS0DK002YM07"
/cell_type="HELA CELLS COT 25-NORMALIZED"
/cell_line="HELA"
/clone_lib="Homo sapiens HELA CELLS COT 25-NORMALIZED"
/note="1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-strand cDNA was digested with Not I and cloned into the Not I and EcoR V sites of the pCMVSPORT 6 vector. Library was normalized."
ORIGIN

Query Match 82.6%; Score 908.4; DB 1; Length 932;
Best Local Similarity 98.6%; Pred. No. 2e-217;
Matches 910; Conservative 5; Mismatches 8; Indels 0; Gaps 0;

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Qy         84 CGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCGACGGGTCATCACGTCGCGCATCG 143
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Db        932 CGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCGACGGGTCATCACTTCGCGCATCG 873

Qy        144 TGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTGGCAGGGGAGCCTGCGCCTGTGGG 203
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Db        872 TGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTGGCAGGGGAGCCTGCGCCTGTGGG 813

Qy        204 ATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGCTGGGCACTCACGGCGGCGCACT 263
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Db        812 ATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGCTGGGAACTCACGGCGGCGCACT 753

Qy        264 GCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGCCAGC 323
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Db        752 GCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGCCAGC 693

Qy        324 TGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTACTACACCCGTTACTTCGTATCGA 383
              |:|||||||||||||||||||||||||||||||||||||||||||::||||||||||||
Db        692 TRACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTACTACACCCGTTWMTTCGTATCGA 633

Qy        384 ATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACCCTATGACATTGCCTTGGTGAAGC 443
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Db        632 ATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACCCTATGACATTGCCTTGGTGAAGC 573

Qy        444 TGTCTGCACCTGTCACCTACACTAAACACATCCAGCCCATCTGTCTCCAGGCCTCCACAT 503
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Db        572 TGTCTGCACCTGTCACCTACACTAAACACATCCAGCCCATCTGTCTCCAGGCCTCCACAT 513

Qy        504 TTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCTGGGGGTACATCAAAGAGGATG 563
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Db        512 TTGAGTTTGAGAACCGGACAAACTGCTGGGTAACTGGCTGGGGGTACATCAAAGAGGATG 453

Qy        564 AGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCAGGTCGCCATCATAAACAACTCTA 623
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Db        452 AGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCAGGTCGCCATCATAAACAACTCTA 393

Qy        624 TGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGGACATCTTTGGAGACATGGTTT 683
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Db        392 TGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGGACATCTTTGGAGACATGGTTT 333

Qy        684 GTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTCGGTGACTCAGGTGGACCCTTGG 743
              |||||||||||||||||||||||||||||||||||| ||||||||||||||||||||||
Db        332 GTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTCGGTAACTCAGGTGGACCCTTGG 273

Qy        744 CCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGTCGTGAGCTGGGGAGTGGGCTGTG 803
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Db        272 CCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGTCGTGAGCTGGGGAGTGGGCTGTG 213

Qy        804 GTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGCCACCACTTTGAGTGGATCCAGA 863
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        212 GTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGCCACCACTTTGAGTGGATCCAGA 153

Qy        864 AGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCCCTCCTGGCCGCTACTCTTTTTCC 923
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        152 AGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCCCTCCTGGCCGCTACTCTTTTTCC 93

Qy        924 CTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTGAGCCTACCTGAGCCCATGCAGCC 983
              ||||||:|||||||||||:||||||||||||||||||||||||||||||||||||||||
Db         92 CTCTTCHCTGGGCTCTCCCMCTCCTGGGGCCGGTCTGAGCCTACCTGAGCCCATGCAGCC 33

Qy        984 TGGGGCCACTGCCAAGTCAGGCC 1006
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                    |||||||||||||||||  ||| ||
Db         32 TGGGGCCACTGCCAAGNCAGCCC 10


RESULT 4
AL555870
LOCUS       AL555870                884 bp    mRNA    linear   EST 30-MAR-2004
DEFINITION  AL555870 Homo sapiens HELA CELLS COT 25-NORMALIZED Homo sapiens
            cDNA clone CS0DK002YM07 5-PRIME, mRNA sequence.
ACCESSION   AL555870
VERSION     AL555870.3  GI:45860593
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 884)
  AUTHORS   Li,W.B., Gruber,C., Jessee,J. and Polayes,D.
  TITLE     Full-length cDNA libraries and normalization
  JOURNAL   Unpublished (2001)
COMMENT     On Feb 15, 2001 this sequence version replaced gi:31277675.
            Contact: Genoscope
            Genoscope - Centre National de Sequencage
            2 rue Gaston Cremieux, CP 5706 - 91057 EVRY cedex - FRANCE
            Email: seqref@genoscope.cns.fr, Web : www.genoscope.cns.fr
            1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime
            end enriched, double-strand cDNA was digested with Not I and cloned
            into the Not I and EcoR V sites of the pCMVSPORT 6 vector. Library
            was normalized. Library was constructed by Life Technologies, a
            division of Invitrogen. This sequence belongs to sequence cluster
            8290.f
            For more information about this cluster, see
            http://www.genoscope.cns.fr/cdna?s=CS0DK002AG04QP1&c=8290.f.
FEATURES             Location/Qualifiers
     source          1. .884
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="CS0DK002YM07"
                     /cell_type="HELA CELLS COT 25-NORMALIZED"
                     /cell_line="HELA"
                     /clone_lib="Homo sapiens HELA CELLS COT 25-NORMALIZED"
                     /note="1st strand cDNA was primed with a NotI-oligo(dT)
                     primer. Five prime end enriched, double-strand cDNA was
                     digested with Not I and cloned into the Not I and EcoR V
                     sites of the pCMVSPORT 6 vector. Library was normalized."
ORIGIN

  Query Match             79.1%;  Score 870.4;  DB 1;  Length 884;
  Best Local Similarity   99.8%;  Pred. No. 7.1e-208;
  Matches  882;  Conservative    0;  Mismatches    1;  Indels    1;  Gaps    1;

Qy          3 CGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCGGG 62
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 CGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCGGG 60

Qy         63 CTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGT-TATCAGGACCATGCGGCCGA 121
              |||||||||||||||||||||||||||||||||||||| ||||||||||||||||||||
Db         61 CTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTGTATCAGGACCATGCGGCCGA 120

Qy        122 CGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTGG 181
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 CGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTGG 180

Qy        182 CAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 CAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGC 240

Qy        242 TGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGGG 301
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 TGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGGG 300

Qy        302 TGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTAC 361
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        301 TGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTAC 360

Qy        362 TACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACCC 421
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        361 TACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACCC 420

Qy        422 TATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCCC 481
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        421 TATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCCC 480

Qy        482 ATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGC 541
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        481 ATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGC 540

Qy        542 TGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCAG 601
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        541 TGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCAG 600

Qy        602 GTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAG 661
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        601 GTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAG 660

Qy        662 GACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTC 721
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        661 GACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTC 720

Qy        722 GGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGTC 781
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        721 GGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGTC 780

Qy        782 GTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGC 841
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        781 GTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGC 840

Qy        842 CACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCAT 885
              |||||||||||||||||||||||||||||||||||||||||||
Db        841 CACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCAT 884


RESULT 5
BX417595
LOCUS       BX417595                1057 bp    mRNA    linear   EST 01-MAY-2004
DEFINITION  BX417595 Homo sapiens PLACENTA Homo sapiens cDNA clone CS0DE010YP05
            5-PRIME, mRNA sequence.
ACCESSION   BX417595
VERSION     BX417595.2  GI:46933208
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 1057)
  AUTHORS   Li,W.B., Gruber,C., Jessee,J. and Polayes,D.
  TITLE     Full-length cDNA libraries and normalization
  JOURNAL   Unpublished (2001)
COMMENT     On May 13, 2003 this sequence version replaced gi:30654423.
            Contact: Genoscope
            Genoscope - Centre National de Sequencage
            2 rue Gaston Cremieux, CP 5706 - 91057 EVRY cedex - FRANCE
            Email: seqref@genoscope.cns.fr, Web : www.genoscope.cns.fr
            1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime
            end enriched, double-strand cDNA was digested with Not I and cloned
            into the Not I and EcoRV sites of the pCMVSPORT 6 vector. Library
            was not normalized. Library was constructed by Life Technologies, a
            division of Invitrogen.
            This sequence belongs to sequence cluster 8290.f
            For more information about this cluster, see
            http://www.genoscope.cns.fr/cdna?s=CS0DE010CH03QP1&c=8290.f.
FEATURES             Location/Qualifiers
     source          1. .1057
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
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/db_xref="taxon:9606"
/clone="CS0DE010YP05"
/tissue_type="PLACENTA"
/clone_lib="Homo sapiens PLACENTA"
/note="Vector: pCMVSPORT_6; 1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-strand cDNA was digested with Not I and cloned into the Not I and EcoRV sites of the pCMVSPORT 6 vector. Library was not normalized."

ORIGIN

Query Match 78.7%; Score 866; DB 5; Length 1057;
Best Local Similarity 95.3%; Pred. No. 9.4e-207;
Matches 942; Conservative 13; Mismatches 23; Indels 10; Gaps 6;

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Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              |||||||||||||||||||||||||||||||||| :|||||||||||||||||||||||||
Db          3 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGC-GTGCTGCTGGCGCTGCTGCTGGCTCG 61

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGT-TATCAGGACCATGCGGCC 119
              |||||||||||||||||||||||||||||||||||||||||| ||||||||||||||||||
Db         62 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTNTATCAGGACCATGCGGCC 121

Qy        120 GACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGT 179
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        122 GACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGT 181

Qy        180 GGCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACC 239
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 GGCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACC 241

Qy        240 GCTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCG 299
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        242 GCTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCG 301

Qy        300 GGTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCT 359
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        302 GGTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCT 361

Qy        360 ACTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCAC 419
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        362 ACTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCAC 421

Qy        420 CCTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGC 479
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        422 CCTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGC 481

Qy        480 CCATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTG 539
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        482 CCATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTG 541

Qy        540 GCTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTC 599
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        542 GCTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTC 601

Qy        600 AGGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCA 659
              |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||||
Db        602 AGGTCGCCATCATAAACAACTCTATGTGC-ACCACCTCTTCCTCAAGTACAGTTTCCGCA 660

Qy        660 AGGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCT 719
              | ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        661 A-GACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCT 719

Qy        720 TCGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAG 779
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        720 TCGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAG 779

Qy        780 TCGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCA 839
              |:|||||||||||||||||||||||||||||||||||||||||||||||||||||:|||||||
Db        780 TYGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACMAATATCA 839

Qy        840 GCCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACC 899
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        840 GCCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACC 899

Qy        900 CCTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCT 959
              | :| :|:|      |:|||||||||||||:|||| |||||     || :|   | :
Db        900 CTYCTKGSC--GCTATYTTTTTCCCTCTTCTYTGGGGTCTCC----ACTCTKGGGCGGGY 953

Qy        960 GAGCCTACCTGAGCCCATGCAGCCTGGG 987
                ||: |||||||:|||||:|  |||
Db        954 TGAGCTWACTGAGCCMATGCASCTGGGG 981
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RESULT 6
BX454526/c
LOCUS BX454526 898 bp mRNA linear EST 05-MAY-2004
DEFINITION BX454526 Homo sapiens PLACENTA Homo sapiens cDNA clone CS0DE010YP05 3-PRIME, mRNA sequence.
ACCESSION BX454526
VERSION BX454526.1 GI:31019139
KEYWORDS EST.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 898)
AUTHORS Li,W.B., Gruber,C., Jessee,J. and Polayes,D.
TITLE Full-length cDNA libraries and normalization
JOURNAL Unpublished (2001)
COMMENT Contact: Genoscope
Genoscope - Centre National de Sequencage
2 rue Gaston Cremieux, CP 5706 - 91057 EVRY cedex - FRANCE
Email: seqref@genoscope.cns.fr, Web : www.genoscope.cns.fr
1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-strand cDNA was digested with Not I and cloned into the Not I and EcoRV sites of the pCMVSPORT 6 vector. Library was not normalized. Library was constructed by Life Technologies, a division of Invitrogen.
This sequence belongs to sequence cluster 8290.f
For more information about this cluster, see
http://www.genoscope.cns.fr/cdna?s=CS0BAK026CB01NM1&c=8290.f.
FEATURES Location/Qualifiers
source 1. .898
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="CS0DE010YP05"
/tissue_type="PLACENTA"
/clone_lib="Homo sapiens PLACENTA"
/note="Vector: pCMVSPORT_6; 1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-strand cDNA was digested with Not I and cloned into the Not I and EcoRV sites of the pCMVSPORT 6 vector. Library was not normalized."

ORIGIN

Query Match 71.0%; Score 780.6; DB 5; Length 898;
Best Local Similarity 96.5%; Pred. No. 2.8e-185;
Matches 862; Conservative 0; Mismatches 24; Indels 7; Gaps 6;

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Qy        184 GGGGAGCCTGCGCCTGTGGGATT--CCCACGTATGCGGAGTGA-GCCTGCTCAGCCACCG 240
              ||  | || |||||| |||||||  ||| |||||||| ||||| || ||||||| | |||
Db        893 GGAAACCCGGCGCCTTTGGGATTTCCCCCCGTATGCGAAGTGAGGCTTGCTCAGGCCCCG 834

Qy        241 CT-GGGCACTCACGGCGG-CGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCC 298
              | |||||||| |||||| |||  |||||||||||||||||||||| ||||||||||||
Db        833 TTGGGGCACTCCCGGCGGCCGCCTTGCTTTGAAACCTATAGTGACCTAAGTGATCCCTCC 774

Qy        299 GGGTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCC 358
              |||||||||||||||||||| |||||||||||||||||||| |   ||||||||||||||
Db        773 GGGTGGATGGTCCAGTTTGGGCAGCTGACTTCCATGCCATCTTCTGGGAGCCTGCAGGCC 714

Qy        359 -TACTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTC 417
               ||||||||||||  ||||||||||||||||||||||||||||||||||||||||||||
Db        713 TTACTACACCCGTTCTTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTC 654
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Qy        418 A-CCCTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCC 476
              | ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        653 ACCCCTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCC 594

Qy        477 AGCCCATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGA 536
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        593 AGCCCATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGA 534

Qy        537 CTGGCTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAG 596
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        533 CTGGCTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAG 474

Qy        597 TTCAGGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCC 656
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        473 TTCAGGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCC 414

Qy        657 GCAAGGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCT 716
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        413 GCAAGGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCT 354

Qy        717 GCTTCGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTG 776
              |||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||
Db        353 GCTTCGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTG 294

Qy        777 GAGTCGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATA 836
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        293 GAGTCGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATA 234

Qy        837 TCAGCCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAG 896
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        233 TCAGCCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAG 174

Qy        897 ACCCCTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGG 956
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        173 ACCCCTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGG 114

Qy        957 TCTGAGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCT 1016
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        113 TCTGAGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCT 54

Qy       1017 TCTGTCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTT 1069
              ||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db         53 TCTGTCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCT 1
```

RESULT 7
BI334059

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LOCUS       BI334059                871 bp    mRNA    linear   EST 30-JUL-2001
DEFINITION  602997320F1 NIH_MGC_12 Homo sapiens cDNA clone IMAGE:5139509 5',
            mRNA sequence.
ACCESSION   BI334059
VERSION     BI334059.1  GI:15018703
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 871)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: ATCC
             cDNA Library Preparation: Life Technologies, Inc.
             cDNA Library Arrayed by: Incyte Genomics, Inc.
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLAM11343  row: h  column: 06
            High quality sequence stop: 826.
FEATURES             Location/Qualifiers
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                     /clone="IMAGE:5139509"
                     /tissue_type="cervical carcinoma cell line"
                     /lab_host="DH10B"
                     /clone_lib="NIH_MGC_12"
                     /note="Organ: cervix; Vector: pCMV-SPORT6; Site_1: NotI;
                     Site_2: SalI; Cloned unidirectionally.  Primer: Oligo dT.
                     Average insert size 1.4 kb. Library prepared by Life
                     Technologies."
ORIGIN
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Query Match 70.7%; Score 777.2; DB 4; Length 871;
Best Local Similarity 97.9%; Pred. No. 2e-184;
Matches 842; Conservative 0; Mismatches 8; Indels 10; Gaps 5;

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Qy        186 GGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGCTGGG 245
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGCTGGG 60

Qy        246 CACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGGGTGGA 305
              |||||||||||||||||||||||||||||      ||||||||||||||||||||||||||
Db         61 CACTCACGGCGGCGCACTGCTTTGAAAC------TGACCTTAGTGATCCCTCCGGGTGGA 114

Qy        306 TGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTACTACA 365
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        115 TGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTACTACA 174

Qy        366 CCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACCCTATG 425
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        175 CCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACCCTATG 234

Qy        426 ACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCCCATCT 485
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        235 ACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCCCATCT 294

Qy        486 GTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCTGGG 545
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        295 GTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCTGGG 354

Qy        546 GGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCAGGTCG 605
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        355 GGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCAGGTCG 414

Qy        606 CCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGGACA 665
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        415 CCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGGACA 474

Qy        666 TCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTCGGTG 725
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        475 TCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTCGGTG 534

Qy        726 ACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGG-AGTCGTG 784
              |||||||||||||||||||||||||||||| |||||||||||||||||||| ||||||
Db        535 ACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGCAGTCGTG 594

Qy        785 AGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGCCAC 844
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        595 AGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGCCAC 654

Qy        845 CACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCCCTCC 904
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        655 CACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCCCTCC 714

Qy        905 TGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTGAGCC 964
              |||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        715 TGGCCG-TACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTGAGCC 773

Qy        965 TACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTGTCTT 1024
              || |||||||||||||||||||||||||||| ||||||  | ||||||||||||||| |||
```

```
Db        774 TAACTGAGCCCATGCAGCCTGGGGCCACTG-CAAGTCCAGGCCTGGTTCTCTTCTGACTT 832

Qy       1025 GTTTGGTAATAAACACATTC 1044
              | | |||||||||| ||||||
Db        833 G-TAGGTAATAAAAACATTC 851


RESULT 8
BI259237
LOCUS       BI259237                777 bp    mRNA    linear   EST 17-JUL-2001
DEFINITION  602970095F1 NIH_MGC_12 Homo sapiens cDNA clone IMAGE:5109695 5',
            mRNA sequence.
ACCESSION   BI259237
VERSION     BI259237.1  GI:14816376
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 777)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: ATCC
             cDNA Library Preparation: Life Technologies, Inc.
             cDNA Library Arrayed by: Incyte Genomics, Inc.
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLAM11265  row: m  column: 24
            High quality sequence stop: 777.
FEATURES             Location/Qualifiers
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                     /db_xref="taxon:9606"
                     /clone="IMAGE:5109695"
                     /tissue_type="cervical carcinoma cell line"
                     /lab_host="DH10B"
                     /clone_lib="NIH_MGC_12"
                     /note="Organ: cervix; Vector: pCMV-SPORT6; Site_1: NotI;
                     Site_2: SalI; Cloned unidirectionally.  Primer: Oligo dT.
                     Average insert size 1.4 kb. Library prepared by Life
                     Technologies."
ORIGIN

  Query Match             69.1%;  Score 760.4;  DB 4;  Length 777;
  Best Local Similarity   99.7%;  Pred. No. 3.3e-180;
  Matches  772;  Conservative    0;  Mismatches    1;  Indels    1;  Gaps    1;

Qy         68 CTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCGACGGGTC 127
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 CTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCGACGGGTC 60

Qy        128 ATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTGGCAGGGG 187
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTGGCAGGGG 120

Qy        188 AGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGCTGGGCA 247
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 AGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGCTGGGCA 180

Qy        248 CTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGGGTGGATG 307
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 CTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGGGTGGATG 240

Qy        308 GTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTACTACACC 367
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 GTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTACTACACC 300

Qy        368 CGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACCCTATGAC 427
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        301 CGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACCCTATGAC 360

Qy        428 ATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCCCATCTGT 487
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        361 ATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCCCATCTGT 420

Qy        488 CTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCTGGGGG 547
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        421 CTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCTGGGGG 480

Qy        548 TACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCAGGTCGCC 607
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        481 TACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCAGGTCGCC 540

Qy        608 ATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGGACATC 667
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        541 ATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGGACATC 600

Qy        668 TTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTCGGTGAC 727
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        601 TTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTCGGTGAC 660

Qy        728 TCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGTCGTGAGC 787
              |||||||||||||||||||||||||| ||||||||||||||||||||||||||||||||
Db        661 TCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGTCGTGAGC 720

Qy        788 TGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGC 841
              |||||||||||||||||||| ||||||||||||||||||||||||||||||||
Db        721 TGGGGAGTGGGCTGTGGTCGG-CCAATCGGCCCGGTGTCTACACCAATATCAGC 773


RESULT 9
BI827608
LOCUS       BI827608                878 bp    mRNA    linear   EST 04-OCT-2001
DEFINITION  603073575F1 NIH_MGC_119 Homo sapiens cDNA clone IMAGE:5165562 5',
            mRNA sequence.
ACCESSION   BI827608
VERSION     BI827608.1  GI:15939145
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 878)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Life Technologies, Inc.
             cDNA Library Preparation: Life Technologies, Inc.
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLAM11411  row: e  column: 19
            High quality sequence stop: 828.
FEATURES             Location/Qualifiers
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                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:5165562"
                     /tissue_type="medulla"
                     /lab_host="DH10B"
                     /clone_lib="NIH_MGC_119"
                     /note="Organ: brain; Vector: pCMV-SPORT6; Site_1: NotI;
                     Site_2: EcoRV (destroyed); RNA source normal medulla from
                     anonymous male age 27.  Library  is oligo-dT primed and
                     directionally cloned (EcoRV site is destroyed  upon
```

cloning). Average insert size 1.3 kb, insert size range 0.9-3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 013. Note: this is a NIH_MGC Library."

ORIGIN

Query Match 66.7%; Score 733.6; DB 4; Length 878;
Best Local Similarity 93.7%; Pred. No. 1.9e-173;
Matches 853; Conservative 0; Mismatches 19; Indels 38; Gaps 7;

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Qy         39 TGCTGGCGCTGCTGCTGGCTCGGGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGC 98
Db          1 TGCTGGCGCTGCTGCTGGCTCGGGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGC 60

Qy         99 CGTTATCAGGACCATGCGGCCGACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACG 158
Db         61 CGTTATCAGGACCATGCGGCCGACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACG 120

Qy        159 CCGAACTCGGGCGTTGGCCGTGGCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCG 218
Db        121 CCGAACTCGGGCG-TGGCCGTGGCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCG 179

Qy        219 GAGTGAGCCTGCTCAGCCACCGCTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATA 278
Db        180 GAGTGAGCCTGCTCAGCCACCGCTGGGCACTCACGGCGGCGCACTGCTTTGAAAC----- 234

Qy        279 GTGACCTTAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCAT 338
Db        235 -TGACCTTAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCAT 293

Qy        339 CCTTCTGGAGCCTGCAGGCCTACTACACCCGTTACTTCGTATCGAATATCTATCTGAGCC 398
Db        294 CCTTCTGGAGCCTGCAGGCCTACTACACCCGTTACTTCGTATCGAATATCTATCTGAGCC 353

Qy        399 CTCGCTACCTGGGGAATTCACCCTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCA 458
Db        354 CTCGCTACCTGGGGAATTCACCCTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCA 413

Qy        459 CCTACACTAAACACATCCAGCCCATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACC 518
Db        414 CCTACACTAAACACATCCAGCCCATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACC 473

Qy        519 GGACAGACTGCTGGGTGACTGGCTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTC 578
Db        474 GGACAGACTGCTGGGTGACTGGCTGGGGGTACATCAAAGAGGATGAG------------- 520

Qy        579 CCCACACCCTCCAGGAAGTTCAGGTCGCCATCATAAACAACTCTATGTGCAACCACCTCT 638
Db        521 ------------GGAAGTTCAGGTCGCCATCATAAACAACTCTATGTGCAACCACCTCT 567

Qy        639 TCCTCAAGTACAGTTTCCGCAAGGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCC 698
Db        568 TCCTCAAGTACAGTTTCCGCAAGGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCC 627

Qy        699 AAGGCGGGAAGGATGCCTGCTTCGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGA-- 756
Db        628 AAGGCGGGAAGGATGCCTGCTTCGGTGACTCAGGTGGACCCTTGGCCTTGTAACAAGAAT 687

Qy        757 TGGACTGTGGTATCAGATTGGAGTCGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAA-TC 815
Db        688 TGGACTGTGGTATCAGGATGGAGTCGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATTC 747

Qy        816 GGCCCGGTGTCTACACCAATATCAGCCACCACTTTGAG-TGGATCCAGAAGCTGATGGCC 874
Db        748 GGCCCGGTGTCTACACCAATATCAAGCACCACTTTGAGTTGGATCCAGAAGCTGATGGCC 807

Qy        875 CAGAGTGGCATGTCCCAGCCAG-ACCCCTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTG 933
Db        808 CAGAGTGGCCATGTCCAGCCAGAACCCCTCCTGGCCACTATTCTTTTTCCCTCATCTCTG 867

Qy        934 GGCTCTCCCA 943
Db        868 GGCTCTTCCA 877
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RESULT 10
CB529999/c
LOCUS CB529999 743 bp mRNA linear EST 05-AUG-2004
DEFINITION UI-H-FT2-bji-b-24-0-UI.s1 NCI_CGAP_FT2 Homo sapiens cDNA clone UI-H-FT2-bji-b-24-0-UI 3', mRNA sequence.
ACCESSION CB529999
VERSION CB529999.1 GI:29391246
KEYWORDS EST.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 743)
AUTHORS NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
TITLE National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index
JOURNAL Unpublished (1997)
COMMENT Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. Gary W. Hunninghake, U of I
cDNA Library preparation: Dr. M. Bento Soares, University of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Distribution information can be found at http://genome.uiowa.edu/distribution/cgap.html
Seq primer: M13 FORWARD
POLYA=Yes.
FEATURES Location/Qualifiers
source 1. .743
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="UI-H-FT2-bji-b-24-0-UI"
/tissue_type="Alveolar Macrophage"
/dev_stage="Adult"
/lab_host="DH10B (Life Technologies)"
/clone_lib="NCI_CGAP_FT2"
/note="Organ: Lung; Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site_1: EcoR I; Site_2: Not I; NCI_CGAP_FT2 is a subtracted cDNA library constructed from a pool of 81 RNA samples from Alveolar Macrophages challenged with different treatments. The mRNA samples were a mixture of these conditions (times refer to incubations following isolation by bronchoalveolar lavage) (some normal donor macrophages were cultured in some of the conditions, other donor macrophages in different conditions). The mRNA samples were pooled for library construction. Control 0 hours; control 3 hours; control 24 hours; LPS 100 ng/ml, 3 hours; LPS 100 ng/ml, 24 hours; PMA 10 ng/ml, 3 hours; PMA 10 ng/ml, 24 hours; Klebsiella moi 10, 3 hours; Klebsiella moi 10, 24 hours; Staph aureus moi 10, 3 hours; Staph aureus moi 10, 24 hours; Adenoviral vector (Ad5 CMV eGFP), moi 500, 3 hours; Adenoviral vector (Ad5 CMV eGFP), moi 500, 24 hours; wt adenovirus moi 500, 3 hours; wt adenovirus moi 500, 24 hours; Ad vector + LPS 3 hours; Ad vector + LPS 24 hours; wt adenovirus + LPS 3 hours; wt adenovirus + LPS 24 hours. The library was subtracted according to Bonaldo, Lennon and Soares, Genome Research, 6:791-806, 1996. The tissue was provided by Dr. Gary W. Hunninghake of the University of Iowa.
TAG_TISSUE=Human Lung Alveolar Macrophage
TAG_LIB=UI-H-FT2
TAG_SEQ=GGCCATGCCG"
ORIGIN

Query Match 60.9%; Score 669.8; DB 6; Length 743;
Best Local Similarity 94.1%; Pred. No. 2.1e-157;
Matches 723; Conservative 0; Mismatches 3; Indels 42; Gaps 1;

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Qy        284 CTTAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTC 343
              ||||||||||||||||||||||||||||||| |||||||||||||||||||||||||||||
Db        743 CTTAGTGATCCCTCCGGGTGGATGGTCCAGTNTGGCCAGCTGACTTCCATGCCATCCTTC 684

Qy        344 TGGAGCCTGCAGGCCTACTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGC 403
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        683 TGGAGCCTGCAGGCCTACTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGC 624

Qy        404 TACCTGGGGAATTCACCCTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTAC 463
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        623 TACCTGGGGAATTCACCCTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTAC 564

Qy        464 ACTAAACACATCCAGCCCATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACA 523
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        563 ACTAAACACATCCAGCCCATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACA 504

Qy        524 GACTGCTGGGTGACTGGCTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCAC 583
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Db        503 GACTGCTGGGTGACTGGCTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCAC 444

Qy        584 ACCCTCCAGGAAGTTCAGGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTC 643
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        443 ACCCTCCAGGAAGTTCAGGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTC 384

Qy        644 AAGTACAGTTTCCGCAAGGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGC 703
              ||||||||||||||||||||||||||||||||||
Db        383 AAGTACAGTTTCCGCAAGGACATCTTTGGAGACATG---------------------------- 348

Qy        704 GGGAAGGATGCCTGCTTCGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTG 763
                                  |||||||||||||||||||||||||||||||| ||||||||
Db        347 ------------------GGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTG 306

Qy        764 TGGTATCAGATTGGAGTCGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGT 823
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        305 TGGTATCAGATTGGAGTCGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGT 246

Qy        824 GTCTACACCAATATCAGCCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGC 883
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Db        245 GTCTACACCAATATCAGCCACCACTTTGAGTGGATCCAGATGCTGATGGCCCAGAGTGGC 186

Qy        884 ATGTCCCAGCCAGACCCCTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCA 943
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        185 ATGTCCCAGCCAGACCCCTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCA 126

Qy        944 CTCCTGGGGCCGGTCTGAGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAG 1003
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        125 CTCCTGGGGCCGGTCTGAGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAG 66

Qy       1004 GCCCTGGTTCTCTTCTGTCTTGTTTGGTAATAAACACATTCCAGTTGA 1051
              ||||||||||||||||||||||||||||||||||||||||||||||||
Db         65 GCCCTGGTTCTCTTCTGTCTTGTTTGGTAATAAACACATTCCAGTTGA 18


RESULT 11
AI126185/c
LOCUS       AI126185                633 bp    mRNA    linear   EST 28-OCT-1998
DEFINITION  qd81h05.x1 Soares_testis_NHT Homo sapiens cDNA clone IMAGE:1735929
            3' similar to SW:PSS8_HUMAN Q16651 PROSTASIN PRECURSOR ;, mRNA
            sequence.
ACCESSION   AI126185
VERSION     AI126185.1  GI:3594699
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 633)
  AUTHORS   NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
  TITLE     National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
            Tumor Gene Index
  JOURNAL   Unpublished (1997)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima
            Bonaldo, Ph.D.
             cDNA Library Arrayed by: Greg Lennon, Ph.D.
             DNA Sequencing by: Washington University Genome Sequencing Center
             Clone distribution: NCI-CGAP clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            www-bio.llnl.gov/bbrp/image/image.html
            Insert Length: 912   Std Error: 0.00
            Seq primer: -40m13 fwd. ET from Amersham
            High quality sequence stop: 455.
FEATURES             Location/Qualifiers
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                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:1735929"
                     /sex="male"
                     /lab_host="DH10B"
                     /clone_lib="Soares_testis_NHT"
                     /note="Vector: pT7T3D-Pac (Pharmacia) with a modified
                     polylinker; Site_1: Not I; Site_2: Eco RI; 1st strand cDNA
                     was prepared from mRNA obtained from Clontech
                     Laboratories, Inc., and primed with  a Not I - oligo(dT)
                     primer [5'
                     TGTTACCAATCTGAAGTGGGAGCGGCCGCCCAATTTTTTTTTTTTTTTT 3']. 
                     Double-stranded cDNA was ligated to Eco RI adaptors
                     (Pharmacia), digested with Not I and cloned into the Not I
                     and Eco RI sites of the modified pT7T3 vector. Library
                     went through one round of normalization to Cot5, and was
                     constructed  by Bento Soares and M. Fatima Bonaldo. "
ORIGIN

  Query Match           56.8%;  Score 624.8;  DB 1;  Length 633;
  Best Local Similarity   99.4%;  Pred. No. 4.3e-146;
  Matches  626;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy        425 GACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCCCATC 484
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        633 GACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCCCATC 574

Qy        485 TGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCTGG 544
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        573 TGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCTGG 514

Qy        545 GGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCAGGTC 604
              |||||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||
Db        513 GGGTACATCAAAGAGGATGAGGCACTGCCATCTCNCCACACCCTCCAGGAAGTTCAGGTC 454

Qy        605 GCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGGAC 664
              |||||||||  ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        453 GCCATCATAACNAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGGAC 394

Qy        665 ATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTCGGT 724
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        393 ATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTCGGT 334

Qy        725 GACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGTCGTG 784
              |||||||||||||||||||||||||||||| |||||||||||||||||||||||||||||||
Db        333 GACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGTCGTG 274

Qy        785 AGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGCCAC 844
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        273 AGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGCCAC 214

Qy        845 CACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCCCTCC 904
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        213 CACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCCCTCC 154

Qy        905 TGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTGAGCC 964
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        153 TGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTGAGCC 94

Qy        965 TACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTGTCTT 1024
```

```
Db         93 TACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTGTCTT 34

Qy       1025 GTTTGGTAATAAACACATTCCAGTTGATGC 1054
Db         33 GTTTGGTAATAAACACATTCCAGTTGATGC 4


RESULT 12
BM019026
LOCUS       BM019026                705 bp    mRNA    linear   EST 30-OCT-2001
DEFINITION  603647010F1 NIH_MGC_98 Homo sapiens cDNA clone IMAGE:5428539 5',
            mRNA sequence.
ACCESSION   BM019026
VERSION     BM019026.1  GI:16533380
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 705)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: ATCC
             cDNA Library Preparation: Ling Hong/Rubin Laboratory
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM1896  row: c  column: 04
            High quality sequence start: 7
            High quality sequence stop: 675.
FEATURES             Location/Qualifiers
     source          1. .705
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:5428539"
                     /tissue_type="astrocytoma grade IV, cell line"
                     /lab_host="DH10B (phage-resistant)"
                     /clone_lib="NIH_MGC_98"
                     /note="Organ: brain; Vector: pOTB7; Site_1: XhoI; Site_2:
                     EcoRI; cDNA made by oligo-dT priming.  Directionally
                     cloned into EcoRI/XhoI sites using the following 5'
                     adaptor: GGCACGAG(G).  Library constructed  by Ling Hong
                     in the laboratory of Gerald M. Rubin (University of
                     California, Berkeley) using ZAP-cDNA synthesis kit
                     (Stratagene) and Superscript II RT (Life Technologies).
                     Note: this is a NIH_MGC Library."
ORIGIN

  Query Match           51.7%;  Score 569.2;  DB 4;  Length 705;
  Best Local Similarity   93.1%;  Pred. No. 4.3e-132;
  Matches  650;  Conservative    0;  Mismatches   18;  Indels   30;  Gaps    4;

Qy          27 GCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCGGGCTGGACTCAGGAAGCCGGAGTCGC 86
Db           2 GCTGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCGGGCT-GACTCAGGAGGCCGGAGTCGC 60

Qy          87 AGGAGGCGGCGCCGTTATCAGGACCATGCGGCCGACGGGTCATCACGTCGCGCATCGTGG 146
Db          61 A-GAGGCGGCGCCGTTATCAGGACCATGCGGCCGACGGGTCATCAGGTCGCGCATCGTGG 119

Qy         147 GTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTGGCAGGGGAGCCTGCGCCTGTGGGATT 206
Db         120 GTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTGGCAGGGGAGCCTGCGCCTGTGGGATT 179

Qy         207 CCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGCTGGGCACTCACGGCGGCGCACTGCT 266
Db         180 CCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGCTGGGCACTCACGGCGGCGCACTGCT 239

Qy         267 TTGAAACCTATAGTGACCTTAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGCCAGCTGA 326
Db         240 TTGAAACCTATAGTGACCTTAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGCCAGCTGA 299

Qy         327 CTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTACTACACCCGTTACTTCGTATCGAATA 386
Db         300 CTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTACTACACCCGTTACTTCGTATCGAATA 359

Qy         387 TCTATCTGAGCCCTCGCTACCTGGGGAATTCACCCTATGACATTGCCTTGGTGAAGCTGT 446
Db         360 TCTATCTGAGCCCTCGCTACCTGGGGAATTCACCCTATGACATTGCCTTGGTGAAGCTGT 419

Qy         447 CTGCACCTGTCACCTACACTAAACACATCCAGCCCATCTGTCTCCAGGCCTCCACATTTG 506
Db         420 CTGCACCTGTCACCTACACTAAACACATCCAGCCCATCTGTCTCCAGGCCTCCACATTTG 479

Qy         507 AGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCTGGGGGTACATCAAAGAGGATGAGG 566
Db         480 AGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCTGGGGGTACATCAAAGAGGATGAG- 538

Qy         567 CACTGCCATCTCCCCACACCCTCCAGGAAGTTCAGGTCGCCATCATAAACAACTCTATGT 626
Db         539 ------------------------GGAAGTTCAGGTCGCCATCATAAACAACTCTATGT 573

Qy         627 GCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGGACATCTTTGGAGACATGGTTTGTG 686
Db         574 GCAACAAACTCTTCCTCAAGTACAGTTTCCGGAAGGACATCTTTTGGAGACATGGTTGTG 633

Qy         687 CTGGCAA--TGCCCAAGGCGGGAAGGATGCCTGCTTCG 722
Db         634 CTGGCAAATGCCCCAAGGCGGGAAGGATGGCTGCTTCG 671


RESULT 13
BM017444
LOCUS       BM017444                682 bp    mRNA    linear   EST 30-OCT-2001
DEFINITION  603644544F1 NIH_MGC_98 Homo sapiens cDNA clone IMAGE:5426338 5',
            mRNA sequence.
ACCESSION   BM017444
VERSION     BM017444.1  GI:16531798
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 682)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: ATCC
             cDNA Library Preparation: Ling Hong/Rubin Laboratory
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM1890  row: g  column: 11
            High quality sequence stop: 672.
FEATURES             Location/Qualifiers
     source          1. .682
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:5426338"
                     /tissue_type="astrocytoma grade IV, cell line"
                     /lab_host="DH10B (phage-resistant)"
                     /clone_lib="NIH_MGC_98"
                     /note="Organ: brain; Vector: pOTB7; Site_1: XhoI; Site_2:
                     EcoRI; cDNA made by oligo-dT priming.  Directionally
```

cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library."

ORIGIN

```
  Query Match            50.0%;  Score 550.4;  DB 4;  Length 682;
  Best Local Similarity  98.8%;  Pred. No. 2.3e-127;
  Matches  565;  Conservative    0;  Mismatches    6;  Indels    1;  Gaps    1;

Qy        106 AGGACCATGCGGCCGACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACT 165
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        111 AGGACCATGCGGCCGACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACT 170

Qy        166 CGGGCGTTGGCC-GTGGCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGA 224
              |||||||||||| ||||||||||||||||||||||||||||||||||||||||||||||
Db        171 CGGGCGTTGGCCTGTGGCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGA 230

Qy        225 GCCTGCTCAGCCACCGCTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACC 284
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        231 GCCTGCTCAGCCACCGCTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACC 290

Qy        285 TTAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCT 344
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        291 TTAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCT 350

Qy        345 GGAGCCTGCAGGCCTACTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCT 404
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        351 GGAGCCTGCAGGCCTACTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCT 410

Qy        405 ACCTGGGGAATTCACCCTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACA 464
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        411 ACCTGGGGAATTCACCCTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACA 470

Qy        465 CTAAACACATCCAGCCCATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAG 524
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        471 CTAAACACATCCAGCCCATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAG 530

Qy        525 ACTGCTGGGTGACTGGCTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACA 584
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        531 ACTGCTGGGTGACTGGCTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCAAA 590

Qy        585 CCCTCCAGGAAGTTCAGGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCA 644
              ||||||||||||||||||||||||||| |||||||||||||||||||||||||||||||||
Db        591 GCCTCCAGGAAGTTCAGGTCGCCATCATACACAACTCTATGTGCAACCACCTCTTCCTCA 650

Qy        645 AGTACAGTTTCCGCAAGGACATCTTTGGAGAC 676
              ||||||||||||| |||||||||  ||||||
Db        651 AGTACAGTTTCCGCCAGGACATCTCGGGAGAC 682
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RESULT 14
BM553333

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LOCUS       BM553333               1219 bp    mRNA    linear   EST 20-FEB-2002
DEFINITION  AGENCOURT_6558382 NIH_MGC_119 Homo sapiens cDNA clone IMAGE:5742958
            5', mRNA sequence.
ACCESSION   BM553333
VERSION     BM553333.1  GI:18791973
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 1219)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Life Technologies, Inc.
            cDNA Library Preparation: Life Technologies, Inc.
            cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
            DNA Sequencing by: Agencourt Bioscience Corporation
            Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLAM12761  row: o  column: 23
            High quality sequence stop: 553.
FEATURES             Location/Qualifiers
     source          1. .1219
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:5742958"
                     /tissue_type="medulla"
                     /lab_host="DH10B"
                     /clone_lib="NIH_MGC_119"
                     /note="Organ: brain; Vector: pCMV-SPORT6; Site_1: NotI;
                     Site_2: EcoRV (destroyed); RNA source normal medulla from
                     anonymous male age 27.  Library  is oligo-dT primed and
                     directionally cloned (EcoRV site is destroyed  upon
                     cloning).  Average insert size 1.3 kb, insert size range
                     0.9-3 kb.  Library is normalized and enriched for
                     full-length clones and  was constructed by C. Gruber
                     (Invitrogen).  Research Genetics  tracking code 013. Note:
                     this is a NIH_MGC Library."
ORIGIN
```

```
  Query Match            49.9%;  Score 548.8;  DB 4;  Length 1219;
  Best Local Similarity  91.8%;  Pred. No. 6.5e-127;
  Matches  602;  Conservative    0;  Mismatches   52;  Indels    2;  Gaps    2;

Qy        106 AGGACCATGCGGCCGACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACT 165
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         66 AGGACCATGCGGCCGACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACT 125

Qy        166 CGGGCGTTGGCCGTGGCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAG 225
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        126 CGGGCGTTGGCCGTGGCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAG 185

Qy        226 CCTGCTCAGCCACCGCTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCT 285
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        186 CCTGCTCAGCCACCGCTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCT 245

Qy        286 TAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTG 345
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        246 TAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTG 305

Qy        346 GAGCCTGCAGGCCTACTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTA 405
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        306 GAGCCTGCAGGCCTACTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTA 365

Qy        406 CCTGGGGAATTCACCCTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACAC 465
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        366 CCTGGGGAATTCACCCTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACAC 425

Qy        466 TAAACACATCCAGCCCATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGA 525
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        426 TAAACACATCCAGCCCATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGA 485

Qy        526 CTGCTGGGTGACTGGCTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACAC 585
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        486 CTGCTGGGTGACTGGCTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACAC 545

Qy        586 CCTCCAGGAAGTTCAGGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAA 645
              |||||||| | |||||| |||||||||| |||||||||| |||||||||||||||||||
Db        546 CCTCCAGGGAATTCAGGGCGCCATCATAGACAACTCTATGGGCAACCACCTCTTCCTCAA 605

Qy        646 GTACA-GTTTCCGCAAGGACATC-TTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGC 703
              ||||| ||| |||||||||||| ||||||||| ||||||       | |    |
Db        606 GTACAGGTTCCCGCAAGGACATCTTTTGGAGACTTGGTTTTGTGCTGGCAGATGCCCAAG 665

Qy        704 GGGAAGGATGCCTGCTTCGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGG 759
              | | ||| || || | | || ||||| ||| | || | ||
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Db        666 GCGGCGGAGCGATGGCCTGGCTTCCCGGAGGGGCCCTTTGCCAGCCACTCCCAGGG 721


RESULT 15
AA161187/c
LOCUS       AA161187                591 bp    mRNA    linear   EST 16-DEC-1996
DEFINITION  zo59b06.s1 Stratagene pancreas (#937208) Homo sapiens cDNA clone
            IMAGE:591155 3' similar to TR:E218488 E218488 TRYPTASE ;, mRNA
            sequence.
ACCESSION   AA161187
VERSION     AA161187.1  GI:1735441
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 591)
  AUTHORS   Hillier,L., Lennon,G., Becker,M., Bonaldo,M.F., Chiapelli,B.,
            Chissoe,S., Dietrich,N., DuBuque,T., Favello,A., Gish,W.,
            Hawkins,M., Hultman,M., Kucaba,T., Lacy,M., Le,M., Le,N.,
            Mardis,E., Moore,B., Morris,M., Parsons,J., Prange,C., Rifkin,L.,
            Rohlfing,T., Schellenberg,K., Soares,M.B., Tan,F., Thierry-Meg,J.,
            Trevaskis,E., Underwood,K., Wohldmann,P., Waterston,R., Wilson,R.
            and Marra,M.
  TITLE     Generation and analysis of 280,000 human expressed sequence tags
  JOURNAL   Genome Res. 6 (9), 807-828 (1996)
  MEDLINE   97044478
   PUBMED   8889549
COMMENT     Contact: Wilson RK
            Washington University School of Medicine
            4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
            Tel: 314 286 1800
            Fax: 314 286 1810
            Email: est@watson.wustl.edu
            This clone is available royalty-free through LLNL ; contact the
            IMAGE Consortium (info@image.llnl.gov) for further information.
            Seq primer: -40M13 fwd. from Amersham
            High quality sequence stop: 386.
FEATURES             Location/Qualifiers
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                     /mol_type="mRNA"
                     /db_xref="GDB:4622541"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:591155"
                     /lab_host="SOLR cells (kanamycin resistant)"
                     /clone_lib="Stratagene pancreas (#937208)"
                     /note="Organ: pancreas; Vector: pBluescript SK-; Site_1:
                     EcoRI; Site_2: XhoI; Cloned unidirectionally.  Primer:
                     Oligo dT. Pancreatic adenocarcinoma cell line. Average
                     insert size: 1.0 kb; Uni-ZAP XR Vector; ~5' adaptor
                     sequence: 5' GAATTCGGCACGAG 3' ~3' adaptor sequence: 5'
                     CTCGAGTTTTTTTTTTTTTTTT 3'"
ORIGIN

  Query Match           47.8%;  Score 525.8;  DB 1;  Length 591;
  Best Local Similarity   97.0%;  Pred. No. 3.5e-121;
  Matches  574;  Conservative    0;  Mismatches   14;  Indels    4;  Gaps    4;

Qy        487 TCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCTGGGG 546
              |||||||||||| ||||| |||||||||||||||| |||||||||| ||||||||||
Db        590 TCTCCAGGCCTCNACATTNGAGTTTGAGAACCGGACAAACTGCTGGGTNACTGGCTGGGG 531

Qy        547 GTACATCAAAGAGGATGAGGCACTGCCATCTCCCC-ACACCCTCCAGGAAGTTCAGGTCG 605
              ||||||||| |||||||||||||||||||||||| ||||| |||||||||||||| |||
Db        530 GTACATCAANGAGGATGAGGCACTGCCATCTCCCCAACACCTTCCAGGAAGTTCAGNTCG 471

Qy        606 CCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGGACA 665
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        470 CCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGGACA 411

Qy        666 TCTTTGGAGACATGGTTTGTGCTGGC-AATGCCCAAGGCGGGAAGGATGCCTGCTTCGGT 724
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Db        410 TCTTTGGAGACATGGTTTGTGCTGGCAAATGCCCAAGGCGGGAAGGATGCCTGCTTCGGT 351

Qy        725 GACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGTCGTG 784
              |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        350 GACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGTCGTG 291

Qy        785 AGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGCCAC 844
              |||||||||||||||||||   ||||||||||||||||||||||||||||||||||||||
Db        290 AGCTGGGGAGTGGGCTGTGGNTCGCCCAATCGGCCCGGTGTCTACACCAATATCAGCCAC 231

Qy        845 CACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCCCTCC 904
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Search completed: March  5, 2005, 09:07:32
Job time : 4257.77 secs
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GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: March 5, 2005, 04:37:53 ; Search time 722.188 Seconds
(without alignments)
9038.138 Million cell updates/sec

Title: US-10-040-647-5
Perfect score: 1100
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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 5401638 seqs, 2966923429 residues

Total number of hits satisfying chosen parameters: 10803276

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_NA:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
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2	1090.6	99.1	1100	9	US-09-909-320-256	Sequence 256, App
3	1090.6	99.1	1100	9	US-09-909-088B-256	Sequence 256, App
4	1090.6	99.1	1100	9	US-09-905-291A-256	Sequence 256, App
5	1090.6	99.1	1100	9	US-09-902-853-256	Sequence 256, App
6	1090.6	99.1	1100	9	US-09-907-824-256	Sequence 256, App
7	1090.6	99.1	1100	9	US-09-907-841-256	Sequence 256, App
8	1090.6	99.1	1100	10	US-09-904-011-256	Sequence 256, App
9	1090.6	99.1	1100	10	US-09-903-640-256	Sequence 256, App
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11	1090.6	99.1	1100	10	US-09-906-742-256	Sequence 256, App
12	1090.6	99.1	1100	10	US-09-906-838-256	Sequence 256, App
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15	1090.6	99.1	1100	10	US-09-904-859-256	Sequence 256, App
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40	1090.6	99.1	1100	10	US-09-907-575-256	Sequence 256, App
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42	1090.6	99.1	1100	10	US-09-902-759-256	Sequence 256, App
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45	1090.6	99.1	1100	10	US-09-907-979-256	Sequence 256, App

ALIGNMENTS

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; Sequence 5, Application US/10040647
; Publication No. US20030092154A1
;  GENERAL INFORMATION:
;       APPLICANT: (US only) ANTALIS Toni Marie and HOOPER John David
;       TITLE OF INVENTION: NOVEL MOLECULES
;       NUMBER OF SEQUENCES: 30
;       CORRESPONDENCE ADDRESS:
;            ADDRESSEE: SCULLY, SCOTT, MURPHY & PRESSER
;            STREET: 400 GARDEN CITY PLAZA
;            CITY: GARDEN CITY
;            STATE: NEW YORK
;            COUNTRY: USA
;            ZIP: 11530
;       COMPUTER READABLE FORM:
;            MEDIUM TYPE: Floppy disk
;            COMPUTER: IBM PC compatible
;            OPERATING SYSTEM: PC-DOS/MS-DOS
;            SOFTWARE: PatentIn Release #1.0, Version #1.25
;       CURRENT APPLICATION DATA:
;            APPLICATION NUMBER: US/10/040,647
;            FILING DATE: 07-Jan-2002
;            CLASSIFICATION: <Unknown>
;       PRIOR APPLICATION DATA:
;            APPLICATION NUMBER: 09/023,942
;            FILING DATE: <Unknown>
;            APPLICATION NUMBER: PO5101/97
;            FILING DATE: 13-FEB-1997
;            APPLICATION NUMBER: PP0422/97
;            FILING DATE: 18-NOV-1997
;            APPLICATION NUMBER: International PCT Application
;            FILING DATE: 13-FEB-1998
;       ATTORNEY/AGENT INFORMATION:
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;             NAME: DIGIGLIO,  FRANK S
;             REGISTRATION NUMBER: 31,346
;             REFERENCE/DOCKET NUMBER: 11168
;       TELECOMMUNICATION INFORMATION:
;             TELEPHONE: (516) 742 4343
;             TELEFAX: (516) 742 4366
;             TELEX: 230 901 SANS UR
;   INFORMATION FOR SEQ ID NO: 5:
;       SEQUENCE CHARACTERISTICS:
;             LENGTH: 1100 base pairs
;             TYPE: nucleic acid
;             STRANDEDNESS: single
;             TOPOLOGY: linear
;       MOLECULE TYPE: DNA
;       FEATURE:
;             NAME/KEY:  CDS
;             LOCATION:  17..961
;       SEQUENCE DESCRIPTION: SEQ ID NO: 5:
US-10-040-647-5

  Query Match             100.0%;  Score 1100;  DB 14;  Length 1100;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 1100;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

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RESULT 2
US-09-909-320-256
; Sequence 256, Application US/09909320
; Patent No. US20020132240A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/909,320
;  CURRENT FILING DATE:  2002-01-04
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
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;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-909-320-256

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Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
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Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
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Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
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Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 3
US-09-909-088B-256
; Sequence 256, Application US/09909088B
; Patent No. US20020146709A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/909,088B
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;  CURRENT FILING DATE:  2001-07-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-909-088B-256

  Query Match             99.1%;  Score 1090.6;  DB 9;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 0;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
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Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
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Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
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Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
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Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097
              ||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAA 1100


RESULT 4
US-09-905-291A-256
; Sequence 256, Application US/09905291A
; Patent No. US20020160374A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
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;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/905,291A
;  CURRENT FILING DATE:  2001-07-12
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-905-291A-256

  Query Match            99.1%;  Score 1090.6;  DB 9;  Length 1100;
  Best Local Similarity  99.6%;  Pred. No. 0;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
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Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
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Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
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Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
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Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
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Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
              |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 5
US-09-902-853-256
; Sequence 256, Application US/09902853
; Publication No. US20020192659A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
```

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;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/902,853
;  CURRENT FILING DATE:  2001-07-10
;  PRIOR APPLICATION NUMBER: US/09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
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;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
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;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-902-853-256

  Query Match             99.1%;  Score 1090.6;  DB 9;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 0;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAAA 1100

RESULT 6
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US-09-907-824-256
; Sequence 256, Application US/09907824
; Publication No. US20020197671A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/907,824
;  CURRENT FILING DATE:  2001-07-17
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-907-824-256

  Query Match             99.1%;  Score 1090.6;  DB 9;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 0;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              |||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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                 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
                 ||||||||||||||||||||||||||||||||||||||||||||| ||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAAA 1100
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RESULT 7
US-09-907-841-256
; Sequence 256, Application US/09907841
; Publication No. US20020198366A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/907,841
; CURRENT FILING DATE: 2001-11-20
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
; LENGTH: 1100
; TYPE: DNA
; ORGANISM: Homo Sapien

US-09-907-841-256

Query Match 99.1%; Score 1090.6; DB 9; Length 1100;
Best Local Similarity 99.6%; Pred. No. 0;
Matches 1093; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
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Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023
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Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097
              ||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAA 1100


RESULT 8
US-09-904-011-256
; Sequence 256, Application US/09904011
; Publication No. US20030003530A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/904,011
;  CURRENT FILING DATE:  2001-07-11
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-904-011-256

  Query Match             99.1%;  Score 1090.6;  DB 10;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 0;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||||||||| ||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              ||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
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Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAA 1100


RESULT 9
US-09-903-640-256
; Sequence 256, Application US/09903640
; Publication No. US20030017463A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/903,640
;  CURRENT FILING DATE:  2001-07-11
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-903-640-256

  Query Match             99.1%;  Score 1090.6;  DB 10;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 0;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
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Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
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Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
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Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAA 1100
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US-09-908-093-256
; Sequence 256, Application US/09908093
; Publication No. US20030017498A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/908,093
;  CURRENT FILING DATE:  2001-07-17
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-908-093-256

  Query Match             99.1%;  Score 1090.6;  DB 10;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 0;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
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Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
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Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
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Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
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Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
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Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAA 1100


RESULT 11
US-09-906-742-256
; Sequence 256, Application US/09906742
; Publication No. US20030023054A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,742
;  CURRENT FILING DATE:  2001-07-16
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-906-742-256

  Query Match             99.1%;  Score 1090.6;  DB 10;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 0;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
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                ||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
                |||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
                |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
                |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100
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RESULT 12
US-09-906-838-256
; Sequence 256, Application US/09906838
; Publication No. US20030027143A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,838
;  CURRENT FILING DATE:  2001-07-16
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-906-838-256

  Query Match             99.1%;  Score 1090.6;  DB 10;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 0;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543
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Qy       541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy       601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy       661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
            |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||||
Db       664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy       721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
            |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||||
Db       724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy       781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy       841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy       901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
            |||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy       961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy      1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
            |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db      1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy      1081 AAAAAAAAAAAAAAAA 1097
            ||||||||||||||||
Db      1084 AAAAAAAAAAAAAAAA 1100
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RESULT 13
US-09-907-613-256
; Sequence 256, Application US/09907613
; Publication No. US20030027145A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/907,613
;  CURRENT FILING DATE:  2001-07-17
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
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;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-907-613-256

  Query Match            99.1%;  Score 1090.6;  DB 10;  Length 1100;
  Best Local Similarity  99.6%;  Pred. No. 0;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy         1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy        61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy       121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy       181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy       241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy       301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy       361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
                |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
                |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
                |||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAA 1100


RESULT 14
US-09-907-942-256
; Sequence 256, Application US/09907942
; Publication No. US20030027146A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/907,942
;  CURRENT FILING DATE:  2002-01-22
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-907-942-256

  Query Match            99.1%;  Score 1090.6;  DB 10;  Length 1100;
  Best Local Similarity  99.6%;  Pred. No. 0;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
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Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              |||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAA 1100


RESULT 15
US-09-904-859-256
; Sequence 256, Application US/09904859
; Publication No. US20030036060A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/904,859
;  CURRENT FILING DATE:  2001-07-12
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-904-859-256

  Query Match           99.1%;  Score 1090.6;  DB 10;  Length 1100;
  Best Local Similarity  99.6%;  Pred. No. 0;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120

Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180

Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240

Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

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Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420

Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480

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Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540

Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600

Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660

Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720

Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780

Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840

Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900

Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960

Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020

Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097

Db       1084 AAAAAAAAAAAAAAAA 1100
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Job time : 727.188 secs

GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: March 4, 2005, 22:18:30 ; Search time 205.888 Seconds
(without alignments)
8742.154 Million cell updates/sec

Title: US-10-040-647-5
Perfect score: 1100
Sequence: 1 CGCGGGAGAGGAGGCCATGG.....AAAAAAAAAAAAAAAAAAAA 1100

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 1202784 seqs, 818138359 residues

Total number of hits satisfying chosen parameters: 2405568

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA:*
1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
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6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1100	100.0	1100	4	US-09-023-942A-5	Sequence 5, Appli
2	1090.6	99.1	1100	4	US-09-907-794A-256	Sequence 256, App
3	1090.6	99.1	1100	4	US-09-905-125A-256	Sequence 256, App
4	1090.6	99.1	1100	4	US-09-902-775A-256	Sequence 256, App
5	1090.6	99.1	1100	4	US-09-906-700-256	Sequence 256, App
6	1090.6	99.1	1100	4	US-09-903-603A-256	Sequence 256, App
7	1090.6	99.1	1100	4	US-09-904-920A-256	Sequence 256, App
8	1090.6	99.1	1100	4	US-09-909-064-256	Sequence 256, App
9	1090.6	99.1	1100	4	US-09-905-381A-256	Sequence 256, App
10	1090.6	99.1	1100	4	US-09-906-618-256	Sequence 256, App
11	1078	98.0	1094	4	US-09-023-942A-3	Sequence 3, Appli
12	1051.2	95.6	1081	3	US-09-008-271A-15	Sequence 15, Appl
13	751	68.3	1103	3	US-09-386-642-59	Sequence 59, Appl
14	482.2	43.8	959	4	US-09-023-942A-25	Sequence 25, Appl
15	351.6	32.0	3866	4	US-09-023-942A-27	Sequence 27, Appl
16	168.2	15.3	1110	3	US-09-386-653A-1	Sequence 1, Appli
17	165	15.0	1212	4	US-09-620-312D-431	Sequence 431, App
18	164.8	15.0	980	4	US-09-023-942A-30	Sequence 30, Appl
19	157.4	14.3	1130	3	US-09-386-653A-8	Sequence 8, Appli
20	133	12.1	1165	4	US-09-023-942A-28	Sequence 28, Appl
21	133	12.1	1378	4	US-09-907-794A-262	Sequence 262, App
22	133	12.1	1378	4	US-09-905-125A-262	Sequence 262, App
23	133	12.1	1378	4	US-09-902-775A-262	Sequence 262, App
24	133	12.1	1378	4	US-09-906-700-262	Sequence 262, App
25	133	12.1	1378	4	US-09-903-603A-262	Sequence 262, App
26	133	12.1	1378	4	US-09-904-920A-262	Sequence 262, App
27	133	12.1	1378	4	US-09-909-064-262	Sequence 262, App
28	133	12.1	1378	4	US-09-905-381A-262	Sequence 262, App
29	133	12.1	1378	4	US-09-906-618-262	Sequence 262, App
30	133	12.1	1430	3	US-09-386-629-1	Sequence 1, Appli
31	128.4	11.7	1166	3	US-09-386-629-2	Sequence 2, Appli
32	125.8	11.4	942	4	US-09-636-382A-3	Sequence 3, Appli
33	119.6	10.9	2440	4	US-09-949-016-5210	Sequence 5210, Ap
34	119.6	10.9	2440	4	US-09-949-016-5211	Sequence 5211, Ap
35	119.6	10.9	2440	4	US-09-949-016-5212	Sequence 5212, Ap
36	119.4	10.9	1142	3	US-09-386-642-8	Sequence 8, Appli
37	119.4	10.9	1169	3	US-09-386-642-7	Sequence 7, Appli
38	117	10.6	1108	2	US-09-016-366A-14	Sequence 14, Appl
39	117	10.6	1108	2	US-08-978-404B-20	Sequence 20, Appl
40	115.4	10.5	1137	2	US-09-016-366A-18	Sequence 18, Appl
41	115.4	10.5	1137	2	US-08-978-404B-13	Sequence 13, Appl
42	114.8	10.4	1341	4	US-10-177-661-3	Sequence 3, Appli
43	114.8	10.4	1434	4	US-10-177-661-1	Sequence 1, Appli
44	114.8	10.4	1748	4	US-09-879-792-11	Sequence 11, Appl
45	114	10.4	1615	4	US-09-820-002-1	Sequence 1, Appli

ALIGNMENTS

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RESULT 1
US-09-023-942A-5
; Sequence 5, Application US/09023942A
; Patent No. 6479274
;  GENERAL INFORMATION:
;    APPLICANT:  (US only) ANTALIS Toni Marie and HOOPER John David
;    TITLE OF INVENTION:  NOVEL MOLECULES
;    NUMBER OF SEQUENCES:  30
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  SCULLY, SCOTT, MURPHY & PRESSER
;      STREET:  400 GARDEN CITY PLAZA
;      CITY:  GARDEN CITY
;      STATE:  NEW YORK
;      COUNTRY:  USA
;      ZIP:  11530
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/023,942A
;      FILING DATE:  13-FEB-1998
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PO5101/97
;      FILING DATE:  13-FEB-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PP0422/97
;      FILING DATE:  18-NOV-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  International PCT Application
;      FILING DATE:  13-FEB-1998
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  DIGIGLIO,  FRANK S
;      REGISTRATION NUMBER:  31,346
;      REFERENCE/DOCKET NUMBER:  11168
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (516) 742 4343
;      TELEFAX:  (516) 742 4366
;      TELEX:  230 901 SANS UR
;  INFORMATION FOR SEQ ID NO:  5:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1100 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA
;    FEATURE:
;      NAME/KEY:  CDS
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;      LOCATION:  17..961
US-09-023-942A-5

  Query Match           100.0%;  Score 1100;  DB 4;  Length 1100;
  Best Local Similarity  100.0%;  Pred. No. 6.5e-295;
  Matches 1100;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Db          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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Db         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
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Db        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
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Db        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
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Db        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
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Db        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Db        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
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Db        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Db        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
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Db       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080

Qy       1081 AAAAAAAAAAAAAAAAAAAA 1100
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Db       1081 AAAAAAAAAAAAAAAAAAAA 1100


RESULT 2
US-09-907-794A-256
; Sequence 256, Application US/09907794A
; Patent No. 6635468
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/907,794A
;  CURRENT FILING DATE:  2001-07-17
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
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;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-907-794A-256

  Query Match             99.1%;  Score 1090.6;  DB 4;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 2.7e-292;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
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Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
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Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
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Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
              |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 3
US-09-905-125A-256
; Sequence 256, Application US/09905125A
; Patent No. 6664376
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/905,125A
;  CURRENT FILING DATE:  2001-07-12
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
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;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-905-125A-256

  Query Match             99.1%;  Score 1090.6;  DB 4;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 2.7e-292;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

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Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
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Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||||||||| ||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              ||||||||||||||||||||||||||||||||||||| ||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 4
US-09-902-775A-256
; Sequence 256, Application US/09902775A
; Patent No. 6686451
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/902,775A
;  CURRENT FILING DATE:  2001-07-10
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
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; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
; LENGTH: 1100
; TYPE: DNA
; ORGANISM: Homo Sapien
US-09-902-775A-256

Query Match 99.1%; Score 1090.6; DB 4; Length 1100;
Best Local Similarity 99.6%; Pred. No. 2.7e-292;
Matches 1093; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
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Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
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Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
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Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
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Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
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Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
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Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
              |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100
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RESULT 5
US-09-906-700-256
; Sequence 256, Application US/09906700
; Patent No. 6723535
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey

```
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,700
;  CURRENT FILING DATE:  2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-906-700-256

  Query Match             99.1%;  Score 1090.6;  DB 4;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 2.7e-292;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
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Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
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Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
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Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
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Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
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Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              |||||||| |||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
              |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 6
US-09-903-603A-256
; Sequence 256, Application US/09903603A
; Patent No. 6767995
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
```

; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: GNE.1618P2C12
; CURRENT APPLICATION NUMBER: US/09/903,603A
; CURRENT FILING DATE: 2001-07-11
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
; LENGTH: 1100
; TYPE: DNA
; ORGANISM: Homo Sapien
US-09-903-603A-256

Query Match 99.1%; Score 1090.6; DB 4; Length 1100;
Best Local Similarity 99.6%; Pred. No. 2.7e-292;
Matches 1093; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
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RESULT 7
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; Sequence 256, Application US/09904920A
; Patent No. 6806352
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi

; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/904,920A
; CURRENT FILING DATE: 2001-07-13
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
; LENGTH: 1100
; TYPE: DNA
; ORGANISM: Homo Sapien
US-09-904-920A-256

Query Match 99.1%; Score 1090.6; DB 4; Length 1100;
Best Local Similarity 99.6%; Pred. No. 2.7e-292;
Matches 1093; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

```
Qy       1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
Db       4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy      61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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Qy     181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
Db     184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy     241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
Db     244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy     301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
Db     304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy     361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Qy     421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
Db     424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy     481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
Db     484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy     541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
Db     544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy     601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
Db     604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy     661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Qy     721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
Db     724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy     781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
Db     784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy     841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
Db     844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy     901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
Db     904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

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Db     964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

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Qy    1081 AAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 8
US-09-909-064-256
; Sequence 256, Application US/09909064
; Patent No. 6818449
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/909,064
;  CURRENT FILING DATE:  2001-07-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
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;  PRIOR APPLICATION NUMBER: PCT/US99/21090
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;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
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;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-909-064-256

  Query Match             99.1%;  Score 1090.6;  DB 4;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 2.7e-292;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097
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RESULT 9
US-09-905-381A-256
; Sequence 256, Application US/09905381A
; Patent No. 6818746
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/905,381A
;  CURRENT FILING DATE:  2001-07-13
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-905-381A-256

  Query Match             99.1%;  Score 1090.6;  DB 4;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 2.7e-292;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              ||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783
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Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
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Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAAA 1100
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RESULT 10
US-09-906-618-256
; Sequence 256, Application US/09906618
; Patent No. 6828146
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,618
;  CURRENT FILING DATE:  2001-07-16
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 256
;   LENGTH: 1100
;   TYPE: DNA
;   ORGANISM: Homo Sapien
US-09-906-618-256

  Query Match            99.1%;  Score 1090.6;  DB 4;  Length 1100;
  Best Local Similarity  99.6%;  Pred. No. 2.7e-292;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
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Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
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Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603
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Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
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Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
              |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 11
US-09-023-942A-3
; Sequence 3, Application US/09023942A
; Patent No. 6479274
;  GENERAL INFORMATION:
;    APPLICANT:  (US only) ANTALIS Toni Marie and HOOPER John David
;    TITLE OF INVENTION:  NOVEL MOLECULES
;    NUMBER OF SEQUENCES:  30
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  SCULLY, SCOTT, MURPHY & PRESSER
;      STREET:  400 GARDEN CITY PLAZA
;      CITY:  GARDEN CITY
;      STATE:  NEW YORK
;      COUNTRY:  USA
;      ZIP:  11530
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/023,942A
;      FILING DATE:  13-FEB-1998
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PO5101/97
;      FILING DATE:  13-FEB-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PP0422/97
;      FILING DATE:  18-NOV-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  International PCT Application
;      FILING DATE:  13-FEB-1998
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  DIGIGLIO,  FRANK S
;      REGISTRATION NUMBER:  31,346
;      REFERENCE/DOCKET NUMBER:  11168
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (516) 742 4343
;      TELEFAX:  (516) 742 4366
;      TELEX:  230 901 SANS UR
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1094 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  17..955
US-09-023-942A-3

  Query Match             98.0%;  Score 1078;  DB 4;  Length 1094;
  Best Local Similarity   99.5%;  Pred. No. 8.3e-289;
  Matches 1094;  Conservative    0;  Mismatches    0;  Indels    6;  Gaps    1;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Db          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
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Db        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
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Db        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAAC------TGACCTTAGTGATCCCTCCGG 294

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        295 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 354

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Db        355 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 414

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        415 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 474

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Db        475 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 534

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
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Db        535 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 594

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        595 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 654

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        655 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 714

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        715 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 774
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Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        775 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 834

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        835 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 894

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        895 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 954

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        955 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1014

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1015 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1074

Qy       1081 AAAAAAAAAAAAAAAAAAAA 1100
              ||||||||||||||||||||
Db       1075 AAAAAAAAAAAAAAAAAAAA 1094
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RESULT 12
US-09-008-271A-15
; Sequence 15, Application US/09008271A
; Patent No. 6203979
; GENERAL INFORMATION:
; APPLICANT: Bandman, Olga
; Hillman, Jennifer L.
; Yue, Henry
; Guegler, Karl J.
; Corley, Neil C.
; Tang, Tom Y.
; Shah, Purvi
; TITLE OF INVENTION: HUMAN PROTEASE MOLECULES
; NUMBER OF SEQUENCES: 24
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Incyte Pharmaceuticals, Inc.
; STREET: 3174 Porter Dr.
; CITY: Palo Alto
; STATE: CA
; COUNTRY: USA
; ZIP: 94304
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSEQ for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/008,271A
; FILING DATE: 16-Jan-1998
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: <Unknown>
; FILING DATE: <Unknown>
; ATTORNEY/AGENT INFORMATION:
; NAME: Mohan-Peterson, Sheela
; REGISTRATION NUMBER: 41,201
; REFERENCE/DOCKET NUMBER: PF-0458 US
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 650-855-0555
; TELEFAX: 650-845-4166
; INFORMATION FOR SEQ ID NO: 15:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1081 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; IMMEDIATE SOURCE:
; LIBRARY: PROSTUT03
; CLONE: 789927

; SEQUENCE DESCRIPTION: SEQ ID NO: 15 :
US-09-008-271A-15

Query Match 95.6%; Score 1051.2; DB 3; Length 1081;
Best Local Similarity 99.7%; Pred. No. 2.2e-281;
Matches 1053; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

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Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         24 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 83

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||||||
Db         84 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCCTTATCAGGACCATGCGGCCG 143

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        144 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 203

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        204 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 263

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        264 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 323

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        324 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 383

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        384 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 443

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        444 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 503

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        504 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 563

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        564 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 623

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        624 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 683

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        684 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 743

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              ||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        744 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 803

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        804 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 863

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        864 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 923

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        924 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 983

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db        984 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1043

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCT 1056
              |||||||||||||||||||||||||||||||||||
Db       1044 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCT 1079


RESULT 13
US-09-386-642-59
; Sequence 59, Application US/09386642
; Patent No. 6420157
; GENERAL INFORMATION:
;  APPLICANT: Darrow, Andrew
;  APPLICANT: Qi, Jenson
;  APPLICANT: Andrade-Gordon, Patricia
;  TITLE OF INVENTION: Zymogen Activation System
;  FILE REFERENCE: ORT-1028
;  CURRENT APPLICATION NUMBER: US/09/386,642
;  CURRENT FILING DATE: 1999-08-31
;  NUMBER OF SEQ ID NOS: 60
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 59
;   LENGTH: 1103
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence: Nucleic acid
;   OTHER INFORMATION: sequence of human protease F in CFEK2 zymogen
;   OTHER INFORMATION: vector
US-09-386-642-59

  Query Match             68.3%;  Score 751;  DB 3;  Length 1103;
  Best Local Similarity   93.4%;  Pred. No. 4.4e-198;
  Matches  784;  Conservative    0;  Mismatches   55;  Indels   0;  Gaps    0;

Qy        140 ATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTGGCAGGGGAGCCTGCGCCTG 199
              ||||| || ||  | |     ||||||||||||||||||||||||||||||||||||||
Db        139 ATCGTTGGGGGCTATGCTCTAGAACTCGGGCGTTGGCCGTGGCAGGGGAGCCTGCGCCTG 198

Qy        200 TGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGCTGGGCACTCACGGCGGCG 259
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        199 TGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCGCTGGGCACTCACGGCGGCG 258

Qy        260 CACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGC 319
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        259 CACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGC 318

Qy        320 CAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTACTACACCCGTTACTTCGTA 379
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db        319 CAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTACTACAACCGTTACTTCGTA 378

Qy        380 TCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACCCTATGACATTGCCTTGGTG 439
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        379 TCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACCCTATGACATTGCCTTGGTG 438

Qy        440 AAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCCCATCTGTCTCCAGGCCTCC 499
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        439 AAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCCCATCTGTCTCCAGGCCTCC 498

Qy        500 ACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCTGGGGGTACATCAAAGAG 559
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        499 ACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGGCTGGGGGTACATCAAAGAG 558

Qy        560 GATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCAGGTCGCCATCATAAACAAC 619
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        559 GATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCAGGTCGCCATCATAAACAAC 618

Qy        620 TCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGGACATCTTTGGAGACATG 679
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        619 TCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAAGGACATCTTTGGAGACATG 678

Qy        680 GTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTCGGTGACTCAGGTGGACCC 739
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        679 GTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTTCGGTGACTCAGGTGGACCC 738

Qy        740 TTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGTCGTGAGCTGGGGAGTGGGC 799
              ||||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||
Db        739 TTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGTCGTGAGCTGGGGAGTGGGC 798

Qy        800 TGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGCCACCACTTTGAGTGGATC 859
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        799 TGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAGCCACCACTTTGAGTGGATC 858

Qy        860 CAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCCCTCCTGGCCGCTACTCTTT 919
              |||||||||||||||||||||||||||||||||||||||||||||||||| |   ||
Db        859 CAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCCCTCCTGGTCTAGACATCAC 918

Qy        920 TTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTGAGCCTACCTGAGCCCATG 978
              | | || || |   | | | | | |   ||  | |  || | |    ||||
Db        919 CATCACCATCACTAGCGGCCGCTTCCCTTTAGTGAGGGTTAATGCTTCGAGCAGACATG 977


RESULT 14
US-09-023-942A-25
; Sequence 25, Application US/09023942A
; Patent No. 6479274
;  GENERAL INFORMATION:
;    APPLICANT:  (US only) ANTALIS Toni Marie and HOOPER John David
;    TITLE OF INVENTION:  NOVEL MOLECULES
;    NUMBER OF SEQUENCES:  30
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  SCULLY, SCOTT, MURPHY & PRESSER
;      STREET:  400 GARDEN CITY PLAZA
;      CITY:  GARDEN CITY
;      STATE:  NEW YORK
;      COUNTRY:  USA
;      ZIP:  11530
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/023,942A
;      FILING DATE:  13-FEB-1998
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PO5101/97
;      FILING DATE:  13-FEB-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PP0422/97
;      FILING DATE:  18-NOV-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  International PCT Application
;      FILING DATE:  13-FEB-1998
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  DIGIGLIO,  FRANK S
;      REGISTRATION NUMBER:  31,346
;      REFERENCE/DOCKET NUMBER:  11168
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (516) 742 4343
;      TELEFAX:  (516) 742 4366
;      TELEX:  230 901 SANS UR
;  INFORMATION FOR SEQ ID NO:  25:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  959 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  2..856
US-09-023-942A-25

  Query Match             43.8%;  Score 482.2;  DB 4;  Length 959;
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  Best Local Similarity   71.3%;  Pred. No. 1.5e-123;
  Matches  672;  Conservative    0;  Mismatches  253;  Indels   18;  Gaps    2;

Qy        101 TTATCAGGACCATGCGGCCGACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCC 160
              || ||||| || ||||| |    ||  |||| | || || || |||||||| || || ||
Db          8 TTGTCAGGGCCCTGCGGTCACAGGACCATCCCTTCCCGTATAGTGGGTGGCGATGATGCT 67

Qy        161 GAACTCGGGCGTTGGCCGTGGCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGA 220
              || || || || |||||||||||| |||||||||||   | ||||    |||| |||| ||
Db         68 GAGCTTGGCCGCTGGCCGTGGCAAGGGAGCCTGCGTGTATGGGGCAACCACTTATGTGGC 127

Qy        221 GTGAGCCTGCTCAGCCACCGCTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGT 280
              |  | | |||||| || ||||||||   || || || || ||||||||  |||
Db        128 GCAACCTTGCTCAACCGCCGCTGGGTGCTTACAGCTGCCCACTGCTTCCAAA-------- 179

Qy        281 GACCTTAGTGATCCCTCCGGGTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCC 340
              |   ||  ||||| |  |  ||||  |||||||||||  |||||||||||| ||||||
Db        180 -AGGATAACGATCCTTTTGACTGGACAGTCCAGTTTGGTGAGCTGACTTCCAGGCCATCT 238

Qy        341 TTCTGGAGCCTGCAGGCCTACTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCT 400
              ||||||  ||| |||||||| | || |||||||    ||    |||| |  ||||||||
Db        239 CTCTGGAACCTACAGGCCTATTCCAACCGTTACCAAATAGAAGATATTTTCCTGAGCCCC 298

Qy        401 CGCTACCTGGGGAATTCACCCTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACC 460
              |||  || | |  ||| ||||||| ||| || ||||||||||  | || ||||||
Db        299 AAGTACTCGGAGCAGTATCCCAATGACATAGCCCTGCTGAAGCTGTCATCTCCAGTCACC 358

Qy        461 TACACTAAACACATCCAGCCCATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGG 520
              |||| |||   |||||||||||||| |||| |  |||||  |   ||||||||||||
Db        359 TACAATAACTTCATCCAGCCCATCTGCCTCCTGAACTCCACGTACAAGTTTGAGAACCGA 418

Qy        521 ACAGACTGCTGGGTGACTGGCTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCC 580
              || |||||||||||||| |||||||||    ||   ||| ||||||   ||||||||||||
Db        419 ACTGACTGCTGGGTGACCGGCTGGGGGGCTATTGGAGAAGATGAGAGTCTGCCATCTCCC 478

Qy        581 CACACCCTCCAGGAAGTTCAGGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTC 640
              ||||  |||||||||| ||||| || || || || ||||||   |||||  |||||  | | |
Db        479 AACACTCTCCAGGAAGTGCAGGTAGCTATTATCAACAACAGCATGTGTAACCATATGTAC 538

Qy        641 CTCAAGTACAGTTTCCGCAAGGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAA 700
              |||     |||||||  | ||||||  |||||||||||||| |||||||  | | ||
Db        539 AAAAAGCCAGACTTCCGCACGAACATCTGGGGAGACATGGTTTGCGCTGGCACTCCTGAA 598

Qy        701 GGCGGGAAGGATGCCTGCTTCGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGA 760
              || || |||||||||||||| |||||||| || ||||||||||||||   || |||||
Db        599 GGTGGCAAGGATGCCTGCTTTGGTGACTCGGGAGGACCCTTGGCCTGCGACCAGGATACG 658

Qy        761 CTGTGGTATCAGATTGGAGTCGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCC 820
              |||||||||| ||||||| |||||||||||| | |||||||||| ||||||| ||
Db        659 GTGTGGTATCAGGTTGGAGTTGTGAGCTGGGGAATAGGCTGTGGTCGCCCCAATCGCCCT 718

Qy        821 GGTGTCTACACCAATATCAGCCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGT 880
              || ||||| ||||| ||||| || ||||   | |||||||||      |||  ||  | |
Db        719 GGAGTCTATACCAACATCAGTCATCACTACAACTGGATCCAGTCAACCATGATCCGCAAT 778

Qy        881 GGCATGTCCCAGCCAGACCCCTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTC 940
              ||  ||  |  ||| |||||   |      |||||| ||| |  ||||  |||||||  |
Db        779 GGGCTGCTCAGGCCTGACCCAGTCCCCTTGCTACTGTTTCTTACTCTGGCCTGGGCTTCC 838

Qy        941 CCACTCCTGGGGCCGGTCTGAGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGT 1000
              |  | ||| |||| | ||||||| ||  | |  | |   |||| |          | ||
Db        839 TCTTTGCTGAGGCCTGCCTGAGCCCACACGTGTACGTCACACCTGTG---------AGGT 889

Qy       1001 CAGGCCCTGGTTCTCTTCTGTCTTGTTTGGTAATAAACACATT 1043
              ||||   ||  ||| || | ||||| ||| ||||||   ||
Db         890 CAGGGTGTGTCTCTTTTGTATCTTGCTTGCTAATAAACCTGTT 932


RESULT 15
US-09-023-942A-27
; Sequence 27, Application US/09023942A
; Patent No. 6479274
;  GENERAL INFORMATION:
;    APPLICANT:  (US only) ANTALIS Toni Marie and HOOPER John David
;    TITLE OF INVENTION:  NOVEL MOLECULES
;    NUMBER OF SEQUENCES:  30
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  SCULLY, SCOTT, MURPHY & PRESSER
;      STREET:  400 GARDEN CITY PLAZA
;      CITY:  GARDEN CITY
;      STATE:  NEW YORK
;      COUNTRY:  USA
;      ZIP:  11530
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/023,942A
;      FILING DATE:  13-FEB-1998
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PO5101/97
;      FILING DATE:  13-FEB-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PP0422/97
;      FILING DATE:  18-NOV-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  International PCT Application
;      FILING DATE:  13-FEB-1998
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  DIGIGLIO,  FRANK S
;      REGISTRATION NUMBER:  31,346
;      REFERENCE/DOCKET NUMBER:  11168
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (516) 742 4343
;      TELEFAX:  (516) 742 4366
;      TELEX:  230 901 SANS UR
;  INFORMATION FOR SEQ ID NO:  27:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3866 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA
US-09-023-942A-27

  Query Match             32.0%;  Score 351.6;  DB 4;  Length 3866;
  Best Local Similarity   97.5%;  Pred. No. 5.3e-87;
  Matches  357;  Conservative    0;  Mismatches    9;  Indels    0;  Gaps    0;

Qy        710 GATGCCTGCTTCGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTAT 769
              | |||   | |||||||||||||||||||||||||||||||||||| ||||||||||||||
Db       3336 GCTGCTCCCCAGGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTAT 3395

Qy        770 CAGATTGGAGTCGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTAC 829
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3396 CAGATTGGAGTCGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTAC 3455

Qy        830 ACCAATATCAGCCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCC 889
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3456 ACCAATATCAGCCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCC 3515

Qy        890 CAGCCAGACCCCTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTG 949
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3516 CAGCCAGACCCCTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTG 3575

Qy        950 GGGCCGGTCTGAGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTG 1009
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3576 GGGCCGGTCTGAGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTG 3635

Qy       1010 GTTCTCTTCTGTCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTT 1069
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3636 GTTCTCTTCTGTCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCT 3695
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Qy       1070 TCAAAA 1075
             ||||||
Db       3696 TCAAAA 3701
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Search completed: March 5, 2005, 09:20:35
Job time : 209.888 secs

GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: March 4, 2005, 20:46:39 ; Search time 674.27 Seconds
(without alignments)
9657.430 Million cell updates/sec

Title: US-10-040-647-5
Perfect score: 1100
Sequence: 1 CGCGGGAGAGGAGGCCATGG.....AAAAAAAAAAAAAAAAAAAA 1100

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 4390206 seqs, 2959870667 residues

Total number of hits satisfying chosen parameters: 8780412

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : N_Geneseq_16Dec04:*
1: geneseqn1980s:*
2: geneseqn1990s:*
3: geneseqn2000s:*
4: geneseqn2001as:*
5: geneseqn2001bs:*
6: geneseqn2002as:*
7: geneseqn2002bs:*
8: geneseqn2003as:*
9: geneseqn2003bs:*
10: geneseqn2003cs:*
11: geneseqn2003ds:*
12: geneseqn2004as:*
13: geneseqn2004bs:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1098.4	99.9	1100	2	AAV59119	Aav59119 Nucleotid
2	1090.6	99.1	1100	2	AAX52259	Aax52259 Protein P
3	1090.6	99.1	1100	3	ADC78568	Adc78568 Human PRO
4	1090.6	99.1	1100	4	AAF72417	Aaf72417 Human PRO
5	1090.6	99.1	1100	8	ACA60211	Aca60211 Human cDN
6	1090.6	99.1	1100	8	ACD07611	Acd07611 Novel hum
7	1090.6	99.1	1100	8	ABX71659	Abx71659 Human cDN
8	1090.6	99.1	1100	8	ACH06991	Ach06991 Human sec
9	1090.6	99.1	1100	8	ABX96228	Abx96228 Human sec
10	1090.6	99.1	1100	8	ACA05549	Aca05549 cDNA enco
11	1090.6	99.1	1100	8	ACD20216	Acd20216 Human sec
12	1090.6	99.1	1100	8	ACA55019	Aca55019 Novel hum
13	1090.6	99.1	1100	9	ACD19854	Acd19854 Human sec
14	1090.6	99.1	1100	9	ADB29461	Adb29461 Human sec
15	1090.6	99.1	1100	9	ADA18317	Ada18317 Human sec
16	1090.6	99.1	1100	9	ACD67001	Acd67001 Human cDN
17	1090.6	99.1	1100	9	ACD83162	Acd83162 Human PRO
18	1090.6	99.1	1100	9	ADA16292	Ada16292 Human sec
19	1090.6	99.1	1100	9	ADA42437	Ada42437 Human sec
20	1090.6	99.1	1100	9	ACD23340	Acd23340 Human PRO
21	1090.6	99.1	1100	9	ADA16716	Ada16716 Human sec
22	1090.6	99.1	1100	9	ADA13145	Ada13145 Human sec
23	1090.6	99.1	1100	9	ADA42013	Ada42013 Human sec
24	1090.6	99.1	1100	9	ADA17360	Ada17360 Human sec
25	1090.6	99.1	1100	9	ADA42863	Ada42863 Human sec
26	1090.6	99.1	1100	9	ACD23702	Acd23702 Human PRO
27	1090.6	99.1	1100	10	ADB77782	Adb77782 Human sec
28	1090.6	99.1	1100	10	ADB74918	Adb74918 Human sec
29	1090.6	99.1	1100	10	ADC28564	Adc28564 Human sec
30	1090.6	99.1	1100	10	ADC39764	Adc39764 Human sec
31	1090.6	99.1	1100	10	ADC40278	Adc40278 Human sec
32	1090.6	99.1	1100	10	ADC19102	Adc19102 Human sec
33	1090.6	99.1	1100	10	ADC34402	Adc34402 Human sec
34	1090.6	99.1	1100	10	ADC29457	Adc29457 Human sec
35	1090.6	99.1	1100	10	ADC28988	Adc28988 Human sec
36	1090.6	99.1	1100	10	ADC40873	Adc40873 Human sec
37	1090.6	99.1	1100	10	ADC19530	Adc19530 Human sec
38	1090.6	99.1	1100	10	ADC33978	Adc33978 Human sec
39	1090.6	99.1	1100	10	ADC13048	Adc13048 Human sec
40	1090.6	99.1	1100	10	ADC12500	Adc12500 Human sec
41	1090.6	99.1	1100	10	ADD05055	Add05055 Human sec
42	1090.6	99.1	1100	10	ADD04061	Add04061 Human sec
43	1090.6	99.1	1100	10	ADD03637	Add03637 Human sec
44	1090.6	99.1	1100	10	ADE34889	Ade34889 Human sec
45	1090.6	99.1	1100	10	ADD89044	Add89044 Encoding

ALIGNMENTS

RESULT 1
AAV59119
ID AAV59119 standard; DNA; 1100 BP.
XX
AC AAV59119;
XX
DT 07-JAN-1999 (first entry)
XX
DE Nucleotide sequence of long isoform of HELA2.
XX
KW Serine protease; regulation; cell activity; viability; HELA2; ATC2;
KW BCOM3; testisin; fertility; suppressor; testicular germ cell cancer;
KW seminoma; testis-specific expression; antitumour; sperm development;
KW infertility; ss.
XX
OS Homo sapiens.
XX
FH Key Location/Qualifiers
FT CDS 17. .961
FT /*tag= a
FT /product= "HELA2"
XX
PN WO9836054-A1.
XX
PD 20-AUG-1998.
XX
PF 13-FEB-1998; 98WO-AU000085.
XX
PR 13-FEB-1997; 97AU-00005101.
PR 18-NOV-1997; 97AU-00000422.
XX
PA (AMRA-) AMRAD OPERATIONS PTY LTD.
XX
PI Antalis TM, Hooper JD;
XX
DR WPI; 1998-480768/41.
DR P-PSDB; AAW77297.
XX
PT New serine protease(s) and kinase involved in regulating cell activity
PT and viability - particularly the testis-specific protease HELA2 used for
PT modulation of fertility and as tumour suppressor.
XX
PS Claim 6; Page 62-64; 167pp; English.

```
XX
CC   The present sequence represents the nucleotide sequence of the long
CC   isoform of HELA2. cDNA generated from HeLa cells and PAI-2 expressing
CC   HeLa cells was amplified using PCR primers AAV48312-13. Three new
CC   sequences were detected in the 480 bp amplicon. These sequences are
CC   designated HELA2 and ATC2 which have high homology to serine proteases
CC   and BCOM3 which has homology to a kinase. The proteins are involved in or
CC   associated with regulation of cell activity and/or viability.
CC   Administration of recombinant HELA2 (also called testisin) is used to
CC   increase fertility. Downregulation of HELA2 reduces fertility. HELA2 is
CC   also a suppressor of testicular germ cell cancers (seminoma) and is also
CC   expressed in some non-testicular cancers (of colon, pancreas, prostate
CC   and ovary), so is a marker/potential therapeutic target for cancer. The
CC   promoter from the HELA2 gene is useful for testis-specific expression of
CC   other genes, e.g. for gene therapy or modulation of fertility. Drugs that
CC   block activity of HELA2 should have antitumour activity (other than in
CC   testis) while in testis recombinant HELA2 should stop growth of tumours
CC   and normalise sperm development (eliminating the need for orchidectomy).
CC   Identification of mutant forms of HELA2 can be used to diagnose
CC   infertility
XX
SQ   Sequence 1100 BP; 226 A; 319 C; 313 G; 242 T; 0 U; 0 Other;

  Query Match             99.9%;  Score 1098.4;  DB 2;  Length 1100;
  Best Local Similarity   99.9%;  Pred. No. 9e-229;
  Matches 1099;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||||||||||||||||||||||||||||||||||||||||||||||| |||||
Db        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCCGTCTG 960

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080

Qy       1081 AAAAAAAAAAAAAAAAAAAA 1100
              ||||||||||||||||||||
Db       1081 AAAAAAAAAAAAAAAAAAAA 1100


RESULT 2
AAX52259
ID   AAX52259 standard; DNA; 1100 BP.
XX
AC   AAX52259;
XX
DT   25-JUN-1999  (first entry)
XX
DE   Protein PRO303 cDNA clone DNA42551-1217.
XX
KW   Secreted protein; transmembrane protein; human; enterocolitis;
KW   Zollinger-Ellison syndrome; gastrointestinal ulceration;
KW   congenital microvillus atrophy; skin disease; cell growth;
KW   abnormal keratinocyte differentiation; psoriasis; epithelial cancer;
KW   Parkinson's disease; Alzheimer's disease; ALS; neuropathy; fibromodulin;
KW   dermal scarring; Usher Syndrome; Atrophia areata; anti-thrombotic;
KW   wound healing; tissue repair; ss.
XX
OS   Homo sapiens.
XX
PN   WO9914328-A2.
XX
PD   25-MAR-1999.
XX
PF   16-SEP-1998;   98WO-US019330.
XX
PR   17-SEP-1997;   97US-0059113P.
PR   17-SEP-1997;   97US-0059115P.
PR   17-SEP-1997;   97US-0059117P.
PR   17-SEP-1997;   97US-0059119P.
PR   17-SEP-1997;   97US-0059121P.
PR   17-SEP-1997;   97US-0059122P.
PR   17-SEP-1997;   97US-0059184P.
PR   18-SEP-1997;   97US-0059263P.
PR   18-SEP-1997;   97US-0059266P.
PR   15-OCT-1997;   97US-0062125P.
PR   17-OCT-1997;   97US-0062285P.
PR   17-OCT-1997;   97US-0062287P.
PR   21-OCT-1997;   97US-0063486P.
PR   24-OCT-1997;   97US-0062814P.
PR   24-OCT-1997;   97US-0062816P.
PR   24-OCT-1997;   97US-0063045P.
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PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
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PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
XX
PA (GETH) GENENTECH INC.
XX
PI Wood WI, Gurney AL, Goddard A, Pennica D, Chen J, Yuan J;
XX
DR WPI; 1999-229533/19.
DR P-PSDB; AAY13388.
XX
PT New isolated human genes and polypeptides used in, e.g. treatment of
PT gastrointestinal ulceration.
XX
PS Claim 2; Fig 91; 320pp; English.
XX
CC AAX52213-74 encode secreted and transmembrane human proteins, and are
CC obtained from cDNA libraries, prepared from fetal lung, fetal kidney,
CC fetal brain, fetal liver and fetal retina. The encoded polypeptides have
CC specific uses based on their homology to known polypeptides, e.g. PRO211
CC and PRO217 can be used for disorders associated with the preservation and
CC maintenance of gastrointestinal mucosa and the repair of acute and
CC chronic mucosal lesions (e.g. enterocolitis, Zollinger-Ellison syndrome,
CC gastrointestinal ulceration and congenital microvillus atrophy), skin
CC diseases associated with abnormal keratinocyte differentiation (e.g.
CC psoriasis, epithelial cancers such as lung squamous cell carcinoma of the
CC vulva and gliomas), potent effects on cell growth and development,
CC diseases related to growth or survival of nerve cells including
CC Parkinson's disease, Alzheimer's disease, ALS, neuropathies or cancer.
CC PRO265 can be used as for fibromodulin, e.g. for reducing dermal
CC scarring. PRO264 can be used as a target for anti-tumor drugs. PRO533 may
CC be used in the treatment of Usher Syndrome or Atrophia areata; PRO269 can
CC be used as an anti-thrombotic agent; PRO287 polypeptides and portions may
CC have therapeutic applications in wound healing and tissue repair; PRO317
CC can be used for treating problems of the kidney, uterus, endometrium,
CC blood vessels, or related tissue, e.g. in the heart of genital tract
XX
SQ Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

Query Match 99.1%; Score 1090.6; DB 2; Length 1100;
Best Local Similarity 99.6%; Pred. No. 4.4e-227;
Matches 1093; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

```
Qy     1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
Db     4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy    61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
Db    64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy   121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
Db   124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy   181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
Db   184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy   241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
Db   244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy   301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
Db   304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy   361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
Db   364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy   421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
Db   424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy   481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
Db   484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy   541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
Db   544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy   601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
Db   604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy   661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
Db   664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy   721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
Db   724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy   781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
Db   784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy   841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
Db   844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy   901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
Db   904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy   961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
Db   964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy  1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
Db  1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083
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Qy       1081 AAAAAAAAAAAAAAAAA 1097
             |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 3
ADC78568
ID   ADC78568 standard; cDNA; 1100 BP.
XX
AC   ADC78568;
XX
DT   01-JAN-2004  (first entry)
XX
DE   Human PRO303 cDNA.
XX
KW   antiinflamatory; antiulcer; cytostatic; antipsoriatic; antiparkinsonian;
KW   nootropic; neuroprotective; vasotropic; chemotaxic; angiogenic;
KW   neurotrophic; osteopathic; antiasthmatic; antiarthritic; antirheumatic;
KW   antiarteriosclerotic; cardiant; antidiabetic; cerebroprotective;
KW   thrombolytic; immunomodulator; enterocolitis; Zollinger-Ellison syndrome;
KW   gastrointestinal ulceration; psoriasis; cancer; Parkinson's disease;
KW   Alzheimer's; ALS; neuropathy; dermal scarring; wound healing;
KW   nerve repair; thrombosis; bone; cartilage formation; angiogenesis;
KW   asthma; rheumatoid arthritis; multiple sclerosis; inflammatory disorder;
KW   atherosclerosis; cardiac injury; infertility; premature aging; AIDS;
KW   diabetes; stroke; gene therapy; transgenic; PRO; human; ss; gene.
XX
OS   Homo sapiens.
XX
PN   WO200015796-A2.
XX
PD   23-MAR-2000.
XX
PF   15-SEP-1999;   99WO-US021090.
XX
PR   16-SEP-1998;   98WO-US019330.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Chen J,  Goddard A,  Gurney AL,  Hillan K,  Pennica D,  Wood WI;
PI   Yuan J;
XX
DR   WPI; 2000-271434/23.
DR   P-PSDB; ADC78569.
XX
PT   Novel nucleic acids encoding secreted and transmembrane polypeptides with
PT   homology, e.g. to growth and cancer-associated antigens.
XX
PS   Claim 2; SEQ ID NO 256; 355pp; English.
XX
CC   The invention relates to a novel nucleic acid encoding a PRO polypeptide.
CC   The polypeptides and polynucleotides of the invention may be useful as
CC   research tools and as therapeutics for treating enterocolitis, Zollinger-
CC   Ellison syndrome, gastrointestinal ulceration, psoriasis, cancer,
CC   Parkinson's disease, Alzheimer's disease, ALS, neuropathies, dermal
CC   scarring and wound healing, nerve repair, thrombosis, bone and/or
CC   cartilage formation, angiogenesis, asthma, rheumatoid arthritis, multiple
CC   sclerosis, inflammatory disorders, atherosclerosis, cardiac injury,
CC   infertility, premature aging, AIDS, diabetes complications and stroke.
CC   The molecules may also be utilised during gene therapy procedures and
CC   transgenic animal production. The current sequence is that of the human
CC   PRO cDNA of the invention.
XX
SQ   Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

  Query Match             99.1%;  Score 1090.6;  DB 3;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 4.4e-227;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              |||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
              |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100
```

```
RESULT 4
AAF72417
ID   AAF72417 standard; cDNA; 1100 BP.
XX
AC   AAF72417;
XX
DT   24-APR-2001  (first entry)
XX
DE   Human PRO303 cDNA.
XX
KW   Human; PRO; dermatological; antipsoriatic; cytostatic; antiinflammatory;
KW   antiparkinsonian nootropic; neuroprotective; vulnerary; cardiant;
KW   antiangiogenic; vasotropic; antiasthmatic; antirheumatic; cancer;
KW   antiarthritic; antiinfertility; antidiabetic; antiviral; diabetes;
KW   ophthalmological; gene therapy; skin disease; gastrointestinal disorder;
KW   ischaemia; inflammation; ss.
XX
OS   Homo sapiens.
XX
PN   WO200104311-A1.
XX
PD   18-JAN-2001.
XX
PF   22-FEB-2000; 2000WO-US004414.
XX
PR   07-JUL-1999;   99US-0143048P.
PR   26-JUL-1999;   99US-0145698P.
PR   28-JUL-1999;   99US-0146222P.
PR   08-SEP-1999;   99WO-US020594.
PR   13-SEP-1999;   99WO-US020944.
PR   15-SEP-1999;   99WO-US021090.
PR   15-SEP-1999;   99WO-US021547.
PR   05-OCT-1999;   99WO-US023089.
PR   29-NOV-1999;   99WO-US028214.
PR   30-NOV-1999;   99WO-US028313.
PR   02-DEC-1999;   99WO-US028564.
PR   02-DEC-1999;   99WO-US028565.
PR   16-DEC-1999;   99WO-US030095.
PR   20-DEC-1999;   99WO-US030911.
PR   20-DEC-1999;   99WO-US030999.
PR   05-JAN-2000; 2000WO-US000219.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Ashkenazi AJ,  Botstein D,  Desnoyers L,  Eaton DL,  Ferrara N;
PI   Filvaroff E,  Fong S,  Gao W,  Gerber H,  Gerritsen ME,  Goddard A;
PI   Godowski PJ,  Grimaldi CJ,  Gurney AL,  Hillan KJ,  Kljavin IJ;
PI   Mather JP,  Pan J,  Paoni NF,  Roy MA,  Stewart TA,  Tumas D;
PI   Williams PM,  Wood WI;
XX
DR   WPI; 2001-081051/09.
DR   P-PSDB; AAB80256.
XX
PT   Sixty one nucleic acids encoding PRO polypeptides which are useful in the
PT   treatment of skin diseases (e.g. psoriasis), cancers (e.g. lung squamous
PT   cell carcinoma) and neurodegenerative diseases (e.g. Alzheimer's
PT   disease).
XX
PS   Claim 2; Fig 91; 393pp; English.
XX
CC   The present sequence is one of sixty one nucleic acids encoding novel
CC   secreted and transmembrane PRO polypeptides. The PRO polypeptides are
CC   useful for treating skin diseases (e.g. psoriasis), cancers (e.g. lung
CC   squamous cell carcinoma), gastrointestinal disorders (e.g.
CC   enterocolitis), neurodegenerative diseases (e.g. Alzheimer's disease,
CC   Parkinson's disease), wound repair, cardiovascular disorders (e.g.
CC   endometrial bleeding angiogenesis, ischaemias such as coronary ischaemia,
CC   atherosclerosis), inflammatory disorders (e.g. asthma, rheumatoid
CC   arthritis, multiple sclerosis), infertility, AIDS and diabetes and
CC   retinal disorders such as retinitis pigmentosum. The PRO nucleic acids
CC   have applications in molecular biology, including use as hybridization
CC   probes, and in chromosome and gene mapping
XX
SQ   Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

  Query Match            99.1%;  Score 1090.6;  DB 4;  Length 1100;
  Best Local Similarity  99.6%;  Pred. No. 4.4e-227;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              |||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              |||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023
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Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
             ||||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
              |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100
```

RESULT 5
ACA60211
ID ACA60211 standard; cDNA; 1100 BP.
XX
AC ACA60211;
XX
DT 12-JUN-2003 (first entry)
XX
DE Human cDNA for secreted/transmembrane protein PRO303.
XX
KW Human; ss; gene; secreted protein; transmembrane protein; PRO;
KW gene therapy; chromosome identification; chromosome marker.
XX
OS Homo sapiens.
XX
PN US2003003530-A1.
XX
PD 02-JAN-2003.
XX
PF 11-JUL-2001; 2001US-00904011.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 01-DEC-1998; 98WO-US025108.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-329602/31.
DR P-PSDB; ABU71935.
XX
PT New transmembrane polypeptides and nucleic acids encoding the
PT polypeptides, useful in gene therapy, in chromosome identification, as
PT chromosome markers, in generating probes and in tissue typing.
XX
PS Claim 2; Fig 91; 484pp; English.
XX
CC The invention relates to an isolated nucleic acid with at least 80%
CC nucleic acid sequence identity to a nucleotide sequence encoding one of
CC 61 secreted/transmembrane polypeptides, or PRO polypeptides or encoding a
CC PRO protein extracellular domain. Also included are a vector comprising
CC the PRO nucleic acid, a host cell comprising the vector, producing a PRO
CC polypeptide (by culturing the host cell for the expression of the PRO
CC polypeptide, and recovering the PRO polypeptide from the cell culture),
CC an isolated PRO polypeptide (having at least 80% sequence identity to: (
CC a) an amino acid sequence selected from the 61 PRO proteins; (b) an amino
CC acid sequence encoded by a nucleic acid molecule deposited with an ATCC
CC number (detailed in the specification); or (c) an extracellular domain of
CC a PRO polypeptide or to a PRO polypeptide lacking its associated signal
CC peptide), a chimaeric molecule comprising a PRO polypeptide of fused to a
CC heterologous amino acid sequence, an anti-PRO antibody, detecting a
CC PRO245 or PRO1868 in a sample suspected of containing the polypeptide,
CC linking a bioactive molecule to a cell expressing a PRO245 or PRO1868 and
CC modulating at least one biological activity of a cell expressing a PRO245
CC or PRO1868. Nucleic acids which encode PRO can be used to generate either

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CC   transgenic animals or knock-out animals which may be used in the
CC   development and screening of therapeutically useful reagents. The nucleic
CC   acids may also be used in gene therapy, in chromosome identification, as
CC   chromosome markers, or in generating probes. The PRO polypeptides are
CC   useful as molecular markers for protein electrophoresis, and the isolated
CC   nucleic acids may be used for recombinantly expressing those markers. The
CC   PRO polypeptides and nucleic acids may also be used in tissue typing.
CC   Anti-PRO antibodies are useful in diagnostic assays for PRO, and in
CC   affinity purification of PRO from recombinant cell culture or natural
CC   sources. The present sequence encodes a PRO protein
XX
SQ   Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

  Query Match             99.1%;  Score 1090.6;  DB 8;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 4.4e-227;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTGCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG. 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
              |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 6
ACD07611
ID   ACD07611 standard; cDNA; 1100 BP.
XX
AC   ACD07611;
XX
DT   07-AUG-2003  (first entry)
XX
DE   Novel human secreted and transmembrane protein PRO303 cDNA.
XX
KW   Human; secreted and transmembrane protein; PRO; pharmaceutical;
KW   diagnostic; biosensor; bioreactor; Parkinson's disease;
KW   Alzheimer's disease; inflammation; nephritis; wound healing;
KW   nerve repair; collateral blood vessel formation; cancer;
KW   colorectal cancer; haemorrhage; rheumatoid arthritis; diabetes;
KW   cirrhosis; fibrosis; restenosis; dermal fibrotic condition; keloid;
KW   scarring; ischaemia; stroke; hypertension; heart attack; atherosclerosis;
KW   infertility; gene therapy; gene; ss.
XX
OS   Homo sapiens.
XX
PN   US2002197671-A1.
XX
PD   26-DEC-2002.
XX
PF   17-JUL-2001; 2001US-00907824.
XX
PR   17-SEP-1997;   97US-0059113P.
PR   17-SEP-1997;   97US-0059115P.
PR   17-SEP-1997;   97US-0059117P.
PR   17-SEP-1997;   97US-0059119P.
PR   17-SEP-1997;   97US-0059121P.
PR   17-SEP-1997;   97US-0059122P.
PR   17-SEP-1997;   97US-0059184P.
PR   18-SEP-1997;   97US-0059263P.
PR   18-SEP-1997;   97US-0059266P.
PR   15-OCT-1997;   97US-0062125P.
PR   17-OCT-1997;   97US-0062285P.
PR   17-OCT-1997;   97US-0062287P.
PR   21-OCT-1997;   97US-0063486P.
PR   24-OCT-1997;   97US-0062814P.
PR   24-OCT-1997;   97US-0062816P.
PR   24-OCT-1997;   97US-0063045P.
PR   24-OCT-1997;   97US-0063120P.
PR   24-OCT-1997;   97US-0063121P.
PR   24-OCT-1997;   97US-0063127P.
PR   24-OCT-1997;   97US-0063128P.
PR   27-OCT-1997;   97US-0063327P.
PR   27-OCT-1997;   97US-0063329P.
PR   28-OCT-1997;   97US-0063541P.
PR   28-OCT-1997;   97US-0063542P.
PR   28-OCT-1997;   97US-0063544P.
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PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 01-DEC-1998; 98WO-US025108.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-370793/35.
DR P-PSDB; ABO01818.
XX
PT New genes and secreted and transmembrane polypeptides (e.g. PRO245 or
PT PRO335), useful for treating or diagnosing e.g. Alzheimer's disease,
PT cancers, hemorrhage, rheumatoid arthritis, diabetes, cirrhosis, ischemia
PT or strokes.
XX
PS Claim 2; Fig 91; 482pp; English.
XX

CC The invention describes a new isolated nucleic acid molecule comprising
CC the full length coding sequence of the DNA deposited with the American
CC Type Culture Collection (e.g. ATCC Deposit No. 209258) ,or a sequence
CC with at least 80% identity to a DNA encoding a PRO polypeptide comprising
CC any of 61 sequences having 164-1119 amino acids fully defined in the
CC specification. The PRO polypeptides or polynucleotides are useful as
CC pharmaceuticals, diagnostics, biosensors or bioreactors. These are
CC particularly useful for detecting or treating e.g. Parkinson's disease,
CC Alzheimer's disease, inflammations, nephritis, wound healing, nerve
CC repair, collateral blood vessel formation, cancers (e.g. colorectal
CC cancer), haemorrhage (or reduce risk for haemorrhage), rheumatoid
CC arthritis, diabetes, cirrhosis of the liver, fibrosis of the lungs,
CC restenosis, dermal fibrotic conditions (e.g. keloids or scarring),
CC ischaemia, strokes, hypertension, heart attacks, atherosclerosis, or
CC infertility in mammals (e.g. humans, dogs, cats, cattle, horses, sheep,
CC pigs, goats, or rabbits) The PRO polypeptides are useful as targets for
CC therapeutic intervention in these diseases, and diagnostic determination
CC of the presence of these diseases. The PRO polypeptides are also useful
CC as molecular weight markers, or for chromosome identification. The PRO
CC genes are useful as hybridisation probes, or for screening libraries of
CC human cDNA, genomic DNA or mRNA. The PRO genes may also be used in gene
CC therapy, particularly for replacing a defective gene. This sequence
CC encodes a novel human secreted and transmembrane PRO polypeptide
XX
SQ Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

Query Match 99.1%; Score 1090.6; DB 8; Length 1100;
Best Local Similarity 99.6%; Pred. No. 4.4e-227;
Matches 1093; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663
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Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||||| |||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||||| |||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| |||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097
              ||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAA 1100
```

RESULT 7
ABX71659
ID ABX71659 standard; cDNA; 1100 BP.
XX
AC ABX71659;
XX
DT 10-MAR-2003 (first entry)
XX
DE Human cDNA encoding secreted/transmembrane protein PRO303.
XX
KW Human; PRO; secreted protein; transmembrane protein; enterocolitis;
KW gastrointestinal ulceration; skin disease; ss; gene;
KW abnormal keratinocyte differentiation; psoriasis; epithelial cancer;
KW squamous cell carcinoma; Alzheimer's disease; Parkinson's disease;
KW amyotrophic lateral sclerosis; inflammatory disease;
KW rheumatoid arthritis; asthma; multiple sclerosis; organ failure;
KW atherosclerosis; cardiac injury; infertility; birth defect;
KW premature aging; AIDS; acquired immunodeficiency syndrome; cancer;
KW diabetic complication; wound repair.
XX
OS Homo sapiens.
XX
PN US2002132240-A1.
XX
PD 19-SEP-2002.
XX
PF 18-JUL-2001; 2001US-00909320.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 01-DEC-1998; 98WO-US025108.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 06-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;

PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-147434/14.
DR P-PSDB; ABU54391.
XX
PT New PRO polypeptides and nucleic acid molecules, useful in diagnosing or
PT treating inflammatory diseases, organ failure, atherosclerosis, cardiac
PT injury, infertility, cancer, AIDS, Alzheimer's disease or Parkinson's
PT disease.
XX
PS Claim 2; Fig 91; 473pp; English.
XX
CC The invention relates to an isolated PRO polypeptide having at least 80%
CC amino acid sequence identity to: (a) any one of 61 fully defined amino
CC acid sequences given in the specification (appearing as ABU54347-
CC ABU54407); (b) an amino acid sequence encoded by the nucleotide sequence
CC deposited under American Type Culture Collection (accession numbers
CC listed in the specification); (c) any one of the PRO sequences which
CC lacks its associated signal peptide; (d) an extracellular domain of the
CC PRO polypeptide with its associated signal peptide; or (e) an
CC extracellular domain of the PRO polypeptide which lacks its associated
CC signal peptide. Also include are the nucleic acids encoding the PRO
CC polypeptides, vectors, host cells and anti-PRO antibodies. The PRO
CC polypeptides and nucleic acids are useful in diagnosing or treating
CC enterocolitis, gastrointestinal ulceration, skin diseases associated with
CC abnormal keratinocyte differentiation, e.g. psoriasis or epithelial
CC cancers such as squamous cell carcinoma, Alzheimer's disease, Parkinson's
CC disease, amyotrophic lateral sclerosis, inflammatory diseases, e.g.
CC rheumatoid arthritis, asthma or multiple sclerosis, organ failure,
CC atherosclerosis, cardiac injury, infertility, birth defects, premature
CC aging, AIDS, cancer, diabetic complications, or mutations in general. The
CC polypeptides are also useful for wound repair and associated therapies
CC concerned with re-growth of tissue. The nucleotide sequences may be used
CC as hybridisation probes in chromosome and gene mapping, or in generating
CC antisense RNA and DNA. PRO nucleic acids are also useful in preparing PRO
CC polypeptides, in assays to identify other proteins or molecules involved
CC in binding reaction, to generate transgenic animals or knockout animals,
CC which in turn are useful in the development and screening of
CC therapeutically useful reagents, for chromosome identification, and
CC tissue typing. The PRO polypeptides and nucleic acid molecules are also
CC useful in gene therapy, and as molecular weight markers for protein
CC electrophoresis purposes. The anti-PRO antibodies may be used in
CC diagnostic assays for PRO, or for the affinity purification of PRO from
CC recombinant cell culture or natural sources. The present sequence encodes
CC a PRO polypeptide
XX
SQ Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

Query Match 99.1%; Score 1090.6; DB 8; Length 1100;
Best Local Similarity 99.6%; Pred. No. 4.4e-227;
Matches 1093; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              ||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              ||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAAA 1100
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RESULT 8
ACH06991
ID ACH06991 standard; cDNA; 1100 BP.
XX
AC ACH06991;
XX
DT 08-OCT-2003 (first entry)
XX
DE Human secreted/transmembrane polypeptide PRO303 cDNA.
XX
KW Human; gene; ss; abnormal bleeding; gynaecological disease; asthma;
KW hysterectomy; angiogenesis; coronary ischaemic condition; skin disease;
KW gastrointestinal mucosa disorder; acute mucosal lesion; neuropathy; ALS;
KW chronic mucosal lesion; abnormal keratinocyte differentiation; psoriasis;
KW Parkinson's disease; Alzheimer's disease; amyotrophic lateral sclerosis;

KW uncontrolled cell growth; cancer; blood coagulation cascade; thrombosis;
KW haemorrhage; endometrial bleeding; angiogenesis; wound healing; tumour;
KW tissue repair; rheumatoid arthritis; multiple sclerosis; tissue typing.
XX
OS Homo sapiens.
XX
PN US2003044839-A1.
XX
PD 06-MAR-2003.
XX
PF 10-JUL-2001; 2001US-00902903.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
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PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
PR 12-DEC-1997; 97US-0069425P.
PR 04-JUN-1998; 98US-0088026P.
PR 10-SEP-1998; 98US-0099803P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98US-0100262P.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98US-0100858P.
PR 17-SEP-1998; 98WO-US019437.
PR 13-OCT-1998; 98US-0104080P.
PR 20-NOV-1998; 98US-0109304P.
PR 01-DEC-1998; 98WO-US025108.
PR 22-DEC-1998; 98US-0113296P.
PR 07-JUL-1999; 99US-0143048P.
PR 26-JUL-1999; 99US-0145698P.
PR 28-JUL-1999; 99US-0146222P.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-492258/46.
DR P-PSDB; ABO47406.
XX
PT Novel secreted and transmembrane polypeptides and polynucleotides
PT encoding them useful for treating abnormal bleeding involved in
PT gynecological diseases, skin diseases and neurodegenerative diseases.
XX
PS Claim 3; Fig 91; 478pp; English.
XX
CC The invention relates to an isolated PRO polypeptide. PRO317 is useful in
CC diagnosing or treating abnormal bleeding involved in gynecological
CC diseases e.g. to avoid or lessen the need for hysterectomy. PRO317 may
CC also be useful as an agent that affects angiogenesis and PRO317 is useful
CC in anti-tumour indications or in treating coronary ischaemic conditions.
CC PRO211 and PRO217 polypeptides are useful for treating disorders
CC associated with the preservation and maintenance of gastrointestinal
CC mucosa and the repair of acute and chronic mucosal lesions, skin diseases
CC associated with abnormal keratinocyte differentiation (e.g. psoriasis).
CC PRO187 polypeptide is useful for treating Parkinson's disease,
CC Alzheimer's disease, amyotrophic lateral sclerosis (ALS), neuropathies
CC and disease related to uncontrolled cell growth, e.g. cancer. PRO219
CC polypeptide plays a regulatory role in the blood coagulation cascade.
CC PRO246 polypeptides which serves as tumour specific antigens may be
CC exploited as therapeutic targets for anti-tumour drugs. PRO269
CC polypeptide is useful as an antithrombotic agent with reduced risk for
CC haemorrhage as compared with heparin. PRO317 polypeptide is useful in
CC treating endometrial bleeding angiogenesis. PRO287 polypeptides and
CC portion have therapeutic applications in wound healing and tissue repair.
CC PRO234 polypeptides are useful for treating asthma, rheumatoid arthritis,
CC psoriasis and multiple sclerosis. The polypeptide and its nucleic acid
CC are useful for tissue typing. PRO antibodies are useful for
CC immunohistochemical staining and/or assay of sample fluids. Anti-PRO
CC antibodies are useful in diagnostic assays for PRO e.g. detecting its
CC expression in specific cells, tissues or serum and for affinity

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CC   purification of PRO from recombinant cell culture or natural sources. The
CC   present sequence represents cDNA encoding a human secreted/transmembrane
CC   PRO polypeptide
XX
SQ   Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

  Query Match             99.1%;  Score 1090.6;  DB 8;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 4.4e-227;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
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Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
              |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 9
ABX96228
ID   ABX96228 standard; cDNA; 1100 BP.
XX
AC   ABX96228;
XX
DT   13-MAY-2003  (first entry)
XX
DE   Human secreted/transmembrane protein cDNA, #47.
XX
KW   Human; gene; ss; PRO; secreted; transmembrane; pharmaceutical;
KW   diagnostic; biosensor; bioreactor; therapeutic; hyperplasia;
KW   endometriosis; cancer; tumour; ischaemia; coronary arterial disease;
KW   polycystic kidney disease; renal failure; inflammatory response; asthma;
KW   rheumatoid arthritis; psoriasis; multiple sclerosis; gene therapy;
KW   cytostatic; gynecological; cardiant; nephrotropic; hepatotropic;
KW   antiinflammatory.
XX
OS   Homo sapiens.
XX
PN   US2002160374-A1.
XX
PD   31-OCT-2002.
XX
PF   12-JUL-2001; 2001US-00905291.
XX
PR   17-SEP-1997;   97US-0059113P.
PR   17-SEP-1997;   97US-0059115P.
PR   17-SEP-1997;   97US-0059117P.
PR   17-SEP-1997;   97US-0059119P.
PR   17-SEP-1997;   97US-0059121P.
PR   17-SEP-1997;   97US-0059122P.
PR   17-SEP-1997;   97US-0059184P.
PR   18-SEP-1997;   97US-0059263P.
PR   18-SEP-1997;   97US-0059266P.
PR   15-OCT-1997;   97US-0062125P.
PR   17-OCT-1997;   97US-0062285P.
PR   17-OCT-1997;   97US-0062287P.
PR   21-OCT-1997;   97US-0063486P.
PR   24-OCT-1997;   97US-0062814P.
PR   24-OCT-1997;   97US-0062816P.
PR   24-OCT-1997;   97US-0063045P.
PR   24-OCT-1997;   97US-0063120P.
PR   24-OCT-1997;   97US-0063121P.
PR   24-OCT-1997;   97US-0063127P.
PR   24-OCT-1997;   97US-0063128P.
PR   27-OCT-1997;   97US-0063327P.
PR   27-OCT-1997;   97US-0063329P.
PR   28-OCT-1997;   97US-0063541P.
PR   28-OCT-1997;   97US-0063542P.
PR   28-OCT-1997;   97US-0063544P.
PR   28-OCT-1997;   97US-0063549P.
PR   28-OCT-1997;   97US-0063550P.
PR   28-OCT-1997;   97US-0063564P.
PR   29-OCT-1997;   97US-0063435P.
PR   29-OCT-1997;   97US-0063704P.
PR   29-OCT-1997;   97US-0063732P.
PR   29-OCT-1997;   97US-0063734P.
PR   29-OCT-1997;   97US-0063735P.
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PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 01-DEC-1998; 98WO-US025108.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-288105/28.
DR P-PSDB; ABU64543.
XX
PT New secreted and transmembrane PRO polypeptides (e.g. PRO533 or PRO245)
PT and genes encoding them, useful for detecting or treating e.g.
PT hyperplasia, endometriosis, cancers, ischemia, coronary arterial disease
PT or inflammations.
XX
PS Claim 2; Fig 91; 477pp; English.
XX
CC The invention discloses isolated PRO secreted/transmembrane polypeptides
CC and the nucleic acid encoding them. The polypeptides can be used to raise
CC antibodies that specifically bind to the PRO polypeptide, for linking a
CC bioactive molecule to a cell expressing a PRO protein and for modulating
CC at least one biological activity of a cell. The PRO polypeptides or
CC polynucleotides are also useful as pharmaceuticals, diagnostics,
CC biosensors or bioreactors, for detecting or treating e.g. hyperplasia,
CC endometriosis, cancers (e.g. those involving solid tumours), ischaemia,
CC coronary arterial disease, polycystic kidney disease, chronic or acute
CC renal failure, or inflammatory responses (e.g. asthma, rheumatoid
CC arthritis, psoriasis or multiple sclerosis) in mammals. The PRO genes may
CC also be used in gene therapy, particularly for replacing a defective
CC gene. The sequences presented in ABX96017-ABX96378 are the genes
CC encoding, the primers amplifying and the probes detecting the PRO
CC polynucleotides of the invention
XX
SQ Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

Query Match 99.1%; Score 1090.6; DB 8; Length 1100;
Best Local Similarity 99.6%; Pred. No. 4.4e-227;
Matches 1093; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

```
Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903
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Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097
              ||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAA 1100
```

RESULT 10
ACA05549
ID ACA05549 standard; cDNA; 1100 BP.
XX
AC ACA05549;
XX
DT 29-MAY-2003 (first entry)
XX
DE cDNA encoding human secreted protein PRO303.
XX
KW Human; gene therapy; mucosal lesion; ulcer; enterocolitis; skin disease;
KW psoriasis; cancer; lung cancer; colon cancer; nerve cell disease;
KW Alzheimer's disease; Parkinson's disease; Usher syndrome; angiogenesis;
KW atrophia areata; inflammatory disease; asthma; rheumatoid arthritis;
KW ischaemia; ss; gene.
XX
OS Homo sapiens.
XX
PN US2003023054-A1.
XX
PD 30-JAN-2003.
XX
PF 16-JUL-2001; 2001US-00906742.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
PR 12-DEC-1997; 97US-0069425P.
PR 04-JUN-1998; 98US-0088026P.
PR 10-SEP-1998; 98US-0099803P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98US-0100262P.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98US-0100858P.
PR 17-SEP-1998; 98WO-US019437.
PR 13-OCT-1998; 98US-0104080P.
PR 20-NOV-1998; 98US-0109304P.
PR 01-DEC-1998; 98WO-US025108.
PR 22-DEC-1998; 98US-0113296P.
PR 07-JUL-1999; 99US-0143048P.
PR 26-JUL-1999; 99US-0145698P.
PR 28-JUL-1999; 99US-0146222P.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-331485/31.
DR P-PSDB; ABU67389.
XX
PT Sixty one isolated nucleic acids encoding a PRO polypeptide, e.g. PRO245

```
PT   or PRO1868, useful in chromosome and gene mapping, in generating
PT   antisense RNA and DNA, and in treating cancer and Alzheimer's disease.
XX
PS   Example 41; Fig 91; 481pp; English.
XX
CC   The invention relates to sixty one nucleic acids encoding PRO
CC   polypeptides (secreted and transmembrane). The polynucleotide is useful
CC   in molecular biology, including uses as hybridisation probes, in
CC   chromosome and gene mapping, in generating antisense RNA and DNA, and in
CC   gene therapy. The polynucleotide may also be used in preparing PRO
CC   polypeptides by recombinant techniques, and in generating either
CC   transgenic animals or knock-out animals which, in turn, are useful in the
CC   development and screening of therapeutically useful reagents. The PRO
CC   polypeptide or the antibody is used in preparing a medicament for
CC   treating a condition responsive to the polypeptide or antibody, such as
CC   mucosal lesions e.g. ulcers and enterocolitis, skin disease e.g.
CC   psoriasis, cancer e.g. lung cancer and colon cancer, nerve cell disease
CC   e.g. Alzheimer's disease and Parkinson's disease, Usher syndrome,
CC   atrophia areata, angiogenesis, inflammatory disease e.g asthma and
CC   rheumatoid arthritis, ischaemia, and in various diagnostic assays. The
CC   present sequence represents an cDNA which encodes a PRO polypeptide
XX
SQ   Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

  Query Match             99.1%;  Score 1090.6;  DB 8;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 4.4e-227;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              ||||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
              |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 11
ACD20216
ID   ACD20216 standard; cDNA; 1100 BP.
XX
AC   ACD20216;
XX
DT   25-AUG-2003  (first entry)
XX
DE   Human secreted / transmembrane polypeptide PRO303 cDNA.
XX
KW   Human; ss; gene; gene therapy; tumour; tissue typing; obesity; diabetes;
KW   hypoinsulinaemia; hyperinsulinaemia; vascular permeability;
KW   cardiac insufficiency disorder; immune response; regeneration; cartilage;
KW   auditory hair cell; hearing loss; bone disorder; sports injury;
KW   arthritis.
XX
OS   Homo sapiens.
XX
PN   US2003036060-A1.
XX
PD   20-FEB-2003.
XX
PF   12-JUL-2001; 2001US-00904859.
XX
PR   17-SEP-1997;   97US-0059113P.
PR   17-SEP-1997;   97US-0059115P.
PR   17-SEP-1997;   97US-0059117P.
PR   17-SEP-1997;   97US-0059119P.
PR   17-SEP-1997;   97US-0059121P.
PR   17-SEP-1997;   97US-0059122P.
PR   17-SEP-1997;   97US-0059184P.
PR   18-SEP-1997;   97US-0059263P.
PR   18-SEP-1997;   97US-0059266P.
PR   15-OCT-1997;   97US-0062125P.
PR   17-OCT-1997;   97US-0062285P.
PR   17-OCT-1997;   97US-0062287P.
PR   21-OCT-1997;   97US-0063486P.
PR   24-OCT-1997;   97US-0062814P.
PR   24-OCT-1997;   97US-0062816P.
PR   24-OCT-1997;   97US-0063045P.
PR   24-OCT-1997;   97US-0063120P.
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PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
PR 12-DEC-1997; 97US-0069425P.
PR 04-JUN-1998; 98US-0088026P.
PR 10-SEP-1998; 98US-0099803P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98US-0100262P.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98US-0100858P.
PR 17-SEP-1998; 98WO-US019437.
PR 13-OCT-1998; 98US-0104080P.
PR 20-NOV-1998; 98US-0109304P.
PR 01-DEC-1998; 98WO-US025108.
PR 22-DEC-1998; 98US-0113296P.
PR 07-JUL-1999; 99US-0143048P.
PR 26-JUL-1999; 99US-0145698P.
PR 28-JUL-1999; 99US-0146222P.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-417923/39.
DR P-PSDB; ABO14909.
XX
PT Novel secreted and transmembrane polypeptide for modulating biological
PT activity of cell expressing the polypeptide, identifying agonists or
PT antagonists of polypeptide, and as molecular weight markers.
XX
PS Claim 2; Fig 91; 469pp; English.
XX
CC The invention relates to an isolated, secreted and transmembrane
CC polypeptide, termed PRO polypeptide. The polypeptide is useful for
CC identifying agonists or antagonists of the polypeptide, for preparing
CC variants of the polypeptide, as molecular weight markers for protein
CC electrophoresis purpose and the nucleic acid is useful for recombinantly
CC expressing those markers. The polypeptide is also useful as therapeutic
CC agent. PRO is useful in assays to identify other proteins or molecules
CC involved in binding interaction. The nucleic acid is useful as
CC hybridisation probes, in chromosome and gene mapping, in generation of
CC antisense RNA and DNA, in the preparation of PRO polypeptide, for
CC generating transgenic animals or knockout animals which in turn are
CC useful in the development and screening of therapeutically useful
CC reagents, to construct hybridisation probes for mapping the gene which
CC encodes the PRO and for the genetic analysis of individuals with genetic
CC disorders, in gene therapy, for chromosome identification, as chromosome
CC marker, and for generating probes for polymerase chain reaction (PCR),
CC Northern analysis, Southern analysis and Western analysis. PRO antibody
CC is useful in diagnostic assays for PRO, e.g. detecting its expression in
CC specific cells, tissues or serum and for affinity purification of PRO
CC from recombinant cell culture or natural sources. The polypeptide or its
CC antibody is useful for the preparation of medicament for treating
CC conditions which is responsive to the PRO polypeptide or anti-PRO
CC antibody e.g. tumour. The polypeptide and the nucleic acid is useful for
CC tissue typing. The polypeptide is useful for treating obesity, diabetes
CC or hypo- or hyper-insulinaemia and cardiac insufficiency disorders, for
CC inhibiting tumour growth, enhances vascular permeability and immune
CC response, for inducing regeneration of auditory hair cells and for
CC treating hearing loss in mammals and for treating bone and/or cartilage
CC disorders such as sports injuries and arthritis. The present sequence
CC represents cDNA encoding a human secreted and transmembrane PRO
CC polypeptide
XX
SQ Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

Query Match 99.1%; Score 1090.6; DB 8; Length 1100;
Best Local Similarity 99.6%; Pred. No. 4.4e-227;
Matches 1093; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243
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Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097
              ||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAA 1100
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RESULT 12
ACA55019
ID ACA55019 standard; cDNA; 1100 BP.
XX
AC ACA55019;
XX
DT 05-JUN-2003 (first entry)
XX
DE Novel human secreted and transmembrane protein PRO303 cDNA.
XX
KW Human; secreted and transmembrane protein; gene therapy; psoriasis;
KW enterocolitis; gastrointestinal ulceration; skin disease;
KW keratinocyte differentiation; epithelial cancer; Alzheimer's disease;
KW squamous cell carcinoma; Parkinson's disease; inflammatory disease;
KW amyotrophic lateral sclerosis; rheumatoid arthritis; asthma;
KW multiple sclerosis; organ failure; atherosclerosis; cardiac injury;
KW infertility; birth defect; premature aging; AIDS; cancer;
KW diabetic complication; wound repair; tissue re-growth; gene; ss.
XX
OS Homo sapiens.
XX
PN US2003017463-A1.
XX
PD 23-JAN-2003.
XX
PF 11-JUL-2001; 2001US-00903640.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
PR 12-DEC-1997; 97US-0069425P.
PR 04-JUN-1998; 98US-0088026P.
PR 10-SEP-1998; 98US-0099803P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98US-0100262P.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98US-0100858P.

PR 17-SEP-1998; 98WO-US019437.
PR 13-OCT-1998; 98US-0104080P.
PR 20-NOV-1998; 98US-0109304P.
PR 01-DEC-1998; 98WO-US025108.
PR 22-DEC-1998; 98US-0113296P.
PR 07-JUL-1999; 99US-0143048P.
PR 26-JUL-1999; 99US-0145698P.
PR 28-JUL-1999; 99US-0146222P.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-341586/32.
DR P-PSDB; ABU69666.
XX
PT New PRO polypeptides and nucleic acid molecules, useful in diagnosing or
PT treating inflammatory diseases, organ failure, atherosclerosis, cardiac
PT injury, infertility, cancer, AIDS, Alzheimer's disease or Parkinson's
PT disease.
XX
PS Claim 2; Fig 91; 473pp; English.
XX
CC The invention describes sixty one nucleic acids encoding PRO polypeptides
CC (secreted and transmembrane). The PRO polypeptides and nucleic acids are
CC useful in diagnosing or treating enterocolitis, gastrointestinal
CC ulceration, skin diseases associated with abnormal keratinocyte
CC differentiation, e.g. psoriasis or epithelial cancers such as squamous
CC cell carcinoma, Alzheimer's disease, Parkinson's disease, amyotrophic
CC lateral sclerosis, inflammatory diseases, e.g. rheumatoid arthritis,
CC asthma or multiple sclerosis, organ failure, atherosclerosis, cardiac
CC injury, infertility, birth defects, premature aging, AIDS, cancer,
CC diabetic complications, or mutations in general. The polypeptides are
CC also useful for wound repair and associated therapies concerned with re-
CC growth of tissue. The PRO polypeptides and nucleic acid molecules are
CC also useful in gene therapy, and as molecular weight markers for protein
CC electrophoresis purposes. The anti-PRO antibodies may be used in
CC diagnostic assays for PRO, or for the affinity purification of PRO from
CC recombinant cell culture or natural sources. This sequence encodes a
CC novel human PRO polypeptide
XX
SQ Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

Query Match 99.1%; Score 1090.6; DB 8; Length 1100;

Best Local Similarity 99.6%; Pred. No. 4.4e-227;
Matches 1093; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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Qy    1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy   61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
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Db   64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy  121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy  181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy  241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy  301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy  361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy  421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy  481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy  541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy  601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy  661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
          |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db  664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy  721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
          |||||||||||||||||||||||||||||||| |||||||||||||||||||||||||||
Db  724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy  781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy  841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy  901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
          ||||||||| |||||||||||||||||||||||||||||||||||||||||||||||||||
Db  904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy  961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy 1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
          |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
              |||||||||||||||||
Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 13
ACD19854
ID   ACD19854 standard; cDNA; 1100 BP.
XX
AC   ACD19854;
XX
DT   22-AUG-2003  (first entry)
XX
DE   Human secreted / transmembrane polypeptide PRO303 cDNA.
XX
KW   Human; ss; gene; gene therapy; apoptosis; bleeding; tumour; ALS;
KW   gynaecological disease; hysterectomy; angiogenesis; skin disease; cancer;
KW   coronary ischaemic condition; gastrointestinal mucosa disorder; asthma;
KW   mucosal lesion repair; keratinocyte differentiation; psoriasis;
KW   Parkinson's disease; Alzheimer's disease; amyotrophic lateral sclerosis;
KW   neuropathy; blood coagulation cascade disorder; thrombosis; haemorrhage;
KW   neurodegenerative disease; endometrial bleeding; wound healing;
KW   tissue repair; rheumatoid arthritis; multiple sclerosis; tissue typing.
XX
OS   Homo sapiens.
XX
PN   US2003027143-A1.
XX
PD   06-FEB-2003.
XX
PF   16-JUL-2001; 2001US-00906838.
XX
PR   17-SEP-1997;   97US-0059113P.
PR   17-SEP-1997;   97US-0059115P.
PR   17-SEP-1997;   97US-0059117P.
PR   17-SEP-1997;   97US-0059119P.
PR   17-SEP-1997;   97US-0059121P.
PR   17-SEP-1997;   97US-0059122P.
PR   17-SEP-1997;   97US-0059184P.
PR   18-SEP-1997;   97US-0059263P.
PR   18-SEP-1997;   97US-0059266P.
PR   15-OCT-1997;   97US-0062125P.
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PR   21-OCT-1997;   97US-0063486P.
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PR   24-OCT-1997;   97US-0063128P.
PR   27-OCT-1997;   97US-0063327P.
PR   27-OCT-1997;   97US-0063329P.
PR   28-OCT-1997;   97US-0063541P.
PR   28-OCT-1997;   97US-0063542P.
PR   28-OCT-1997;   97US-0063544P.
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PR   28-OCT-1997;   97US-0063550P.
PR   28-OCT-1997;   97US-0063564P.
PR   29-OCT-1997;   97US-0063435P.
PR   29-OCT-1997;   97US-0063704P.
PR   29-OCT-1997;   97US-0063732P.
PR   29-OCT-1997;   97US-0063734P.
PR   29-OCT-1997;   97US-0063735P.
PR   29-OCT-1997;   97US-0063738P.
PR   29-OCT-1997;   97US-0064215P.
PR   31-OCT-1997;   97US-0063870P.
PR   31-OCT-1997;   97US-0064103P.
PR   03-NOV-1997;   97US-0064248P.
PR   07-NOV-1997;   97US-0064809P.
PR   12-NOV-1997;   97US-0065186P.
PR   17-NOV-1997;   97US-0065846P.
PR   18-NOV-1997;   97US-0065693P.
PR   21-NOV-1997;   97US-0066120P.
PR   21-NOV-1997;   97US-0066364P.
PR   24-NOV-1997;   97US-0066453P.
PR   24-NOV-1997;   97US-0066466P.
PR   24-NOV-1997;   97US-0066511P.
PR   24-NOV-1997;   97US-0066770P.
PR   24-NOV-1997;   97US-0066772P.
PR   25-NOV-1997;   97US-0066840P.
PR   12-DEC-1997;   97US-0069425P.
PR   04-JUN-1998;   98US-0088026P.
PR   10-SEP-1998;   98US-0099803P.
PR   10-SEP-1998;   98WO-US018824.
PR   14-SEP-1998;   98US-0100262P.
PR   14-SEP-1998;   98WO-US019177.
PR   16-SEP-1998;   98WO-US019330.
PR   17-SEP-1998;   98US-0100858P.
PR   17-SEP-1998;   98WO-US019437.
PR   13-OCT-1998;   98US-0104080P.
PR   20-NOV-1998;   98US-0109304P.
PR   01-DEC-1998;   98WO-US025108.
PR   22-DEC-1998;   98US-0113296P.
PR   07-JUL-1999;   99US-0143048P.
PR   26-JUL-1999;   99US-0145698P.
PR   28-JUL-1999;   99US-0146222P.
PR   08-SEP-1999;   99WO-US020594.
PR   13-SEP-1999;   99WO-US020944.
PR   15-SEP-1999;   99WO-US021090.
PR   15-SEP-1999;   99WO-US021547.
PR   05-OCT-1999;   99WO-US023089.
PR   29-NOV-1999;   99WO-US028214.
PR   30-NOV-1999;   99WO-US028313.
PR   01-DEC-1999;   99WO-US028301.
PR   02-DEC-1999;   99WO-US028564.
PR   02-DEC-1999;   99WO-US028565.
PR   16-DEC-1999;   99WO-US030095.
PR   20-DEC-1999;   99WO-US030911.
PR   20-DEC-1999;   99WO-US030999.
PR   05-JAN-2000; 2000WO-US000219.
PR   11-FEB-2000; 2000WO-US003565.
PR   22-FEB-2000; 2000WO-US004414.
PR   24-FEB-2000; 2000WO-US005004.
PR   02-MAR-2000; 2000WO-US005841.
PR   20-MAR-2000; 2000WO-US007377.
PR   30-MAR-2000; 2000WO-US008439.
PR   22-MAY-2000; 2000WO-US014042.
PR   02-JUN-2000; 2000WO-US015264.
PR   28-JUL-2000; 2000WO-US020710.
PR   24-AUG-2000; 2000WO-US023328.
PR   18-SEP-2000; 2000US-00665350.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Ashkenazi A,  Botstein D,  Desnoyers L,  Eaton DL,  Ferrara N;
PI   Filvaroff E,  Fong S,  Gao W,  Gerber H,  Gerritsen ME,  Goddard A;
PI   Godowski PJ,  Grimaldi JC,  Gurney AL,  Hillan KJ,  Kljavin IJ;
PI   Mather JP,  Pan J,  Paoni NF,  Roy MA,  Stewart TA,  Tumas D;
PI   Williams PM,  Wood WI;
XX
DR   WPI; 2003-417249/39.
DR   P-PSDB; AB014848.
XX
PT   Novel secreted and transmembrane polypeptides and polynucleotides
PT   encoding them useful for treating abnormal bleeding involved in
PT   gynecological diseases, skin diseases and neurodegenerative diseases.
XX
PS   Claim 2; Fig 91; 467pp; English.
XX
CC   The invention relates to an isolated secreted and transmembrane PRO
CC   polypeptide. The PRO polypeptides are useful for modulating biological
CC   activity of a cell, in diagnosing or treating abnormal bleeding involved
```

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CC   in gynaecological diseases e.g. to avoid or lessen the need for
CC   hysterectomy, for treating angiogenesis, tumour, coronary ischaemic
CC   condition, disorders associated with the preservation and maintenance of
CC   gastrointestinal mucosa and the repair of acute and chronic mucosal
CC   lesions, skin diseases associated with abnormal keratinocyte
CC   differentiation (e.g. psoriasis), Parkinson's disease, Alzheimer's
CC   disease, amyotrophic lateral sclerosis (ALS), neuropathies, disease
CC   related to uncontrolled cell growth (e.g. cancer), blood coagulation
CC   cascade disorders, neurodegenerative disease, thrombosis, haemorrhage,
CC   endometrial bleeding, wound healing, tissue repair, asthma, rheumatoid
CC   arthritis, multiple sclerosis. Nucleic acid encoding PRO polypeptides are
CC   useful in molecular biology including uses as hybridisation probes and in
CC   the generation of antisense RNA and DNA, for preparing PRO polypeptides,
CC   for generating transgenic animals or knockout animals. The PRO
CC   polypeptides and their nucleic acids are useful for tissue typing. PRO
CC   antibodies are useful for immunohistochemical staining and/or assay of
CC   sample fluids. Anti-PRO antibodies are useful in diagnostic assays for
CC   PRO e.g. detecting its expression in specific cells, tissues or serum and
CC   for affinity purification of PRO from recombinant cell culture or natural
CC   sources. The present sequence represents cDNA encoding a human secreted
CC   and transmembrane PRO polypeptide
XX
SQ   Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

  Query Match             99.1%;  Score 1090.6;  DB 9;  Length 1100;
  Best Local Similarity   99.6%;  Pred. No. 4.4e-227;
  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
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Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
              ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
              ||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAA 1100


RESULT 14
ADB29461
ID   ADB29461 standard; cDNA; 1100 BP.
XX
AC   ADB29461;
XX
DT   20-NOV-2003  (first entry)
XX
DE   Human secreted/transmembrane protein cDNA, #49.
XX
KW   Human; gene; ss; PRO; secreted; transmembrane; gastrointestinal mucosa;
KW   mucosal lesion; skin disease; keratinocyte differentiation; psoriasis;
KW   Parkinson's disease; Alzheimer's diseases; amyotrophic lateral sclerosis;
KW   ALS; neuropathy; cell growth; cancer; tumour; viral infection;
KW   neurodegenerative disease; antithrombotic agent; haemorrhage;
KW   endometrial bleeding angiogenesis; kidney tissue; apoptosis; therapeutic;
KW   tissue typing; immunohistochemical staining; gene therapy; nootropic;
KW   neuroprotective; cytostatic; virucide; anticoagulant.
XX
OS   Homo sapiens.
XX
PN   US2003092002-A1.
XX
PD   15-MAY-2003.
XX
PF   10-JUL-2001; 2001US-00902615.
XX
PR   17-SEP-1997;   97US-0059113P.
PR   17-SEP-1997;   97US-0059115P.
PR   17-SEP-1997;   97US-0059117P.
PR   17-SEP-1997;   97US-0059119P.
PR   17-SEP-1997;   97US-0059121P.
PR   17-SEP-1997;   97US-0059122P.
PR   17-SEP-1997;   97US-0059184P.
PR   18-SEP-1997;   97US-0059263P.
PR   18-SEP-1997;   97US-0059266P.
PR   15-OCT-1997;   97US-0062125P.
PR   17-OCT-1997;   97US-0062285P.
PR   17-OCT-1997;   97US-0062287P.
PR   21-OCT-1997;   97US-0063486P.
PR   24-OCT-1997;   97US-0062814P.
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PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
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PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
PR 12-DEC-1997; 97US-0069425P.
PR 04-JUN-1998; 98US-0088026P.
PR 10-SEP-1998; 98US-0099803P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98US-0100262P.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98US-0100858P.
PR 17-SEP-1998; 98WO-US019437.
PR 13-OCT-1998; 98US-0104080P.
PR 20-NOV-1998; 98US-0109304P.
PR 01-DEC-1998; 98WO-US025108.
PR 22-DEC-1998; 98US-0113296P.
PR 07-JUL-1999; 99US-0143048P.
PR 26-JUL-1999; 99US-0145698P.
PR 28-JUL-1999; 99US-0146222P.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005044.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-765473/72.
DR P-PSDB; ADB29462.
XX
PT Novel isolated native PRO polypeptide useful for treating Parkinson's
PT disease, enterocolitis, Zollinger-Ellison syndrome gastrointestinal
PT ulceration, Alzheimer's disease, amyotrophic lateral sclerosis, Usher
PT syndrome.
XX
PS Claim 2; Fig 91; 469pp; English.
XX
CC The invention discloses isolated PRO secreted/transmembrane polypeptides
CC and the nucleic acid encoding them. The polypeptides can be used to raise
CC antibodies that specifically bind to the PRO polypeptide, for linking a
CC bioactive molecule to a cell expressing a PRO protein and for modulating
CC at least one biological activity of a cell. PRO polypeptides are useful
CC for detecting other PRO polypeptides in a sample and for linking a
CC bioactive molecule to a cell expressing a PRO polypeptide. The PRO
CC polypeptide antibodies are useful for modulating the biological activity
CC of a cell expressing PRO polypeptides. PRO polypeptides are also useful
CC for treating disorders associated with the preservation and maintenance
CC of gastrointestinal mucosa and the repair of acute and chronic mucosal
CC lesions, skin diseases associated with abnormal keratinocyte
CC differentiation (e.g. psoriasis), Parkinson's disease, Alzheimer's
CC diseases, amyotrophic lateral sclerosis (ALS), neuropathies and
CC additionally, disease related to uncontrolled cell growth, e.g. cancer.
CC PRO polypeptides also serves as tumour specific antigens which may be
CC exploited as therapeutic targets for anti-tumour drugs, and are also
CC employed therapeutically in vivo for lessening the effects of viral
CC infection. The PRO polypeptides can be also used in assays to determine
CC if it has a role in neurodegenerative diseases or their reversal, as an
CC antithrombotic agent with reduced risk for haemorrhage as compared with
CC heparin, in treating other PRO-associated disorders, in modulating
CC endometrial bleeding angiogenesis, and may also have an effect on kidney
CC tissue. PRO polypeptides and their portions affect the expression of
CC genes which have a role in apoptosis. The polynucleotides are useful in
CC molecular biology including uses as hybridisation probes for cDNA library
CC to isolate the full-length PRO cDNA or to isolate other cDNAs, in
CC chromosome and gene mapping, in the generation of antisense RNA and DNA,
CC for preparing PRO polypeptides, for generating transgenic animals or
CC knockout animals which are useful in the development and screening of
CC therapeutically useful reagents, as probes and for the genetic analysis
CC of individuals with genetic disorders as well as for recombinantly
CC expressing the protein and for chromosome identification. The proteins
CC are useful as molecular marker for protein electrophoresis purposes, as
CC therapeutic agents, for screening compounds to identify those that mimic
CC the PRO polypeptide (agonists) or prevent the effect of the PRO
CC polypeptide (antagonists). The polynucleotides and proteins are useful
CC for tissue typing. PRO antibodies are useful for immunohistochemical
CC staining and/or assay of sample fluids. Anti-PRO antibodies are useful in
CC diagnostic assays for PRO e.g. detecting its expression in specific
CC cells, tissues or serum and for affinity purification of PRO from
CC recombinant cell culture or natural sources. The PRO genes may also be
CC used in gene therapy, particularly for replacing a defective gene. The
CC sequence presented is a gene encoding a PRO polynucleotide of the
CC invention.
XX
SQ Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

Query Match 99.1%; Score 1090.6; DB 9; Length 1100;
Best Local Similarity 99.6%; Pred. No. 4.4e-227;

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  Matches 1093;  Conservative    0;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        304 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 363

Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        364 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 423

Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        424 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 483

Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        484 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 543

Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        544 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 603

Qy        601 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 660
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Db        604 GGTCGCCATCATAAACAACTCTATGTGCAACCACCTCTTCCTCAAGTACAGTTTCCGCAA 663

Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
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Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
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Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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Db        964 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1023

Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAAA 1100


RESULT 15
ADA18317
ID   ADA18317 standard; cDNA; 1100 BP.
XX
AC   ADA18317;
XX
DT   20-NOV-2003  (first entry)
XX
DE   Human secreted/transmembrane protein cDNA, #49.
XX
KW   Human; gene; ss; PRO; secreted; transmembrane; gastrointestinal mucosa;
KW   mucosal lesion; skin disease; keratinocyte differentiation; psoriasis;
KW   Parkinson's disease; Alzheimer's diseases; amyotrophic lateral sclerosis;
KW   ALS; neuropathy; cell growth; cancer; tumour; viral infection;
KW   neurodegenerative disease; antithrombotic agent; haemorrhage;
KW   endometrial bleeding angiogenesis; kidney tissue; apoptosis; therapeutic;
KW   tissue typing; immunohistochemical staining; gene therapy; nootropic;
KW   neuroprotective; cytostatic; virucide; anticoagulant.
XX
OS   Homo sapiens.
XX
PN   US2003039971-A1.
XX
PD   27-FEB-2003.
XX
PF   16-JUL-2001; 2001US-00906646.
XX
PR   17-SEP-1997;   97US-0059113P.
PR   17-SEP-1997;   97US-0059115P.
PR   17-SEP-1997;   97US-0059117P.
PR   17-SEP-1997;   97US-0059119P.
PR   17-SEP-1997;   97US-0059121P.
PR   17-SEP-1997;   97US-0059122P.
PR   17-SEP-1997;   97US-0059184P.
PR   18-SEP-1997;   97US-0059263P.
PR   18-SEP-1997;   97US-0059266P.
PR   15-OCT-1997;   97US-0062125P.
PR   17-OCT-1997;   97US-0062285P.
PR   17-OCT-1997;   97US-0062287P.
PR   21-OCT-1997;   97US-0063486P.
PR   24-OCT-1997;   97US-0062814P.
PR   24-OCT-1997;   97US-0062816P.
PR   24-OCT-1997;   97US-0063045P.
PR   24-OCT-1997;   97US-0063120P.
PR   24-OCT-1997;   97US-0063121P.
PR   24-OCT-1997;   97US-0063127P.
PR   24-OCT-1997;   97US-0063128P.
PR   27-OCT-1997;   97US-0063327P.
PR   27-OCT-1997;   97US-0063329P.
PR   28-OCT-1997;   97US-0063541P.
PR   28-OCT-1997;   97US-0063542P.
PR   28-OCT-1997;   97US-0063544P.
PR   28-OCT-1997;   97US-0063549P.
PR   28-OCT-1997;   97US-0063550P.
PR   28-OCT-1997;   97US-0063564P.
PR   29-OCT-1997;   97US-0063435P.
PR   29-OCT-1997;   97US-0063704P.
PR   29-OCT-1997;   97US-0063732P.
PR   29-OCT-1997;   97US-0063734P.
PR   29-OCT-1997;   97US-0063735P.
PR   29-OCT-1997;   97US-0063738P.
PR   29-OCT-1997;   97US-0064215P.
PR   31-OCT-1997;   97US-0063870P.
PR   31-OCT-1997;   97US-0064103P.
PR   03-NOV-1997;   97US-0064248P.
PR   07-NOV-1997;   97US-0064809P.
PR   12-NOV-1997;   97US-0065186P.
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PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
PR 12-DEC-1997; 97US-0069425P.
PR 04-JUN-1998; 98US-0088026P.
PR 10-SEP-1998; 98US-0099803P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98US-0100262P.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98US-0100858P.
PR 17-SEP-1998; 98WO-US019437.
PR 13-OCT-1998; 98US-0104080P.
PR 20-NOV-1998; 98US-0109304P.
PR 01-DEC-1998; 98WO-US025108.
PR 22-DEC-1998; 98US-0113296P.
PR 07-JUL-1999; 99US-0143048P.
PR 26-JUL-1999; 99US-0145698P.
PR 28-JUL-1999; 99US-0146222P.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-503392/47.
DR P-PSDB; ADA18318.
XX
PT New secreted and transmembrane polypeptides useful for treating skin,
PT neurodegenerative diseases, asthma, rheumatoid arthritis, psoriasis and
PT multiple sclerosis.
XX
PS Claim 2; SEQ ID NO 256; 471pp; English.
XX
CC The invention discloses isolated PRO secreted/transmembrane polypeptides
CC and the nucleic acid encoding them. The polypeptides can be used to raise
CC antibodies that specifically bind to the PRO polypeptide, for linking a
CC bioactive molecule to a cell expressing a PRO protein and for modulating
CC at least one biological activity of a cell. PRO polypeptides are useful
CC for detecting other PRO polypeptides in a sample and for linking a
CC bioactive molecule to a cell expressing a PRO polypeptide. The PRO
CC polypeptide antibodies are useful for modulating the biological activity
CC of a cell expressing PRO polypeptides. PRO polypeptides are also useful
CC for treating disorders associated with the preservation and maintenance
CC of gastrointestinal mucosa and the repair of acute and chronic mucosal
CC lesions, skin diseases associated with abnormal keratinocyte
CC differentiation (e.g. psoriasis), Parkinson's disease, Alzheimer's
CC diseases, amyotrophic lateral sclerosis (ALS), neuropathies and
CC additionally, disease related to uncontrolled cell growth, e.g. cancer.
CC PRO polypeptides also serves as tumour specific antigens which may be
CC exploited as therapeutic targets for anti-tumour drugs, and are also
CC employed therapeutically in vivo for lessening the effects of viral
CC infection. The PRO polypeptides can be also used in assays to determine
CC if it has a role in neurodegenerative diseases or their reversal, as an
CC antithrombotic agent with reduced risk for haemorrhage as compared with
CC heparin, in treating other PRO-associated disorders, in modulating
CC endometrial bleeding angiogenesis, and may also have an effect on kidney
CC tissue. PRO polypeptides and their portions affect the expression of
CC genes which have a role in apoptosis. The polynucleotides are useful in
CC molecular biology including uses as hybridisation probes for cDNA library
CC to isolate the full-length PRO cDNA or to isolate other cDNAs, in
CC chromosome and gene mapping, in the generation of antisense RNA and DNA,
CC for preparing PRO polypeptides, for generating transgenic animals or
CC knockout animals which are useful in the development and screening of
CC therapeutically useful reagents, as probes and for the genetic analysis
CC of individuals with genetic disorders as well as for recombinantly
CC expressing the protein and for chromosome identification. The proteins
CC are useful as molecular marker for protein electrophoresis purposes, as
CC therapeutic agents, for screening compounds to identify those that mimic
CC the PRO polypeptide (agonists) or prevent the effect of the PRO
CC polypeptide (antagonists). The polynucleotides and proteins are useful
CC for tissue typing. PRO antibodies are useful for immunohistochemical
CC staining and/or assay of sample fluids. Anti-PRO antibodies are useful in
CC diagnostic assays for PRO e.g. detecting its expression in specific
CC cells, tissues or serum and for affinity purification of PRO from
CC recombinant cell culture or natural sources. The PRO genes may also be
CC used in gene therapy, particularly for replacing a defective gene. The
CC sequence presented is a gene encoding a PRO polynucleotide of the
CC invention.
XX
SQ Sequence 1100 BP; 225 A; 321 C; 314 G; 240 T; 0 U; 0 Other;

Query Match 99.1%; Score 1090.6; DB 9; Length 1100;
Best Local Similarity 99.6%; Pred. No. 4.4e-227;
Matches 1093; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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Qy          1 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 60
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Db          4 CGCGGGAGAGGAGGCCATGGGCGCGCGCGGGGCGCTGCTGCTGGCGCTGCTGCTGGCTCG 63

Qy         61 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         64 GGCTGGACTCAGGAAGCCGGAGTCGCAGGAGGCGGCGCCGTTATCAGGACCATGCGGCCG 123

Qy        121 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        124 ACGGGTCATCACGTCGCGCATCGTGGGTGGAGAGGACGCCGAACTCGGGCGTTGGCCGTG 183

Qy        181 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        184 GCAGGGGAGCCTGCGCCTGTGGGATTCCCACGTATGCGGAGTGAGCCTGCTCAGCCACCG 243

Qy        241 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        244 CTGGGCACTCACGGCGGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGG 303

Qy        301 GTGGATGGTCCAGTTTGGCCAGCTGACTTCCATGCCATCCTTCTGGAGCCTGCAGGCCTA 360
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Qy        361 CTACACCCGTTACTTCGTATCGAATATCTATCTGAGCCCTCGCTACCTGGGGAATTCACC 420
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Qy        421 CTATGACATTGCCTTGGTGAAGCTGTCTGCACCTGTCACCTACACTAAACACATCCAGCC 480
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Qy        481 CATCTGTCTCCAGGCCTCCACATTTGAGTTTGAGAACCGGACAGACTGCTGGGTGACTGG 540
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Qy        541 CTGGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACCCTCCAGGAAGTTCA 600
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Qy        661 GGACATCTTTGGAGACATGGTTTGTGCTGGCAATGCCCAAGGCGGGAAGGATGCCTGCTT 720
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Db        664 GGACATCTTTGGAGACATGGTTTGTGCTGGCAACGCCCAAGGCGGGAAGGATGCCTGCTT 723

Qy        721 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGGATGGACTGTGGTATCAGATTGGAGT 780
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Db        724 CGGTGACTCAGGTGGACCCTTGGCCTGTAACAAGAATGGACTGTGGTATCAGATTGGAGT 783

Qy        781 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 840
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Db        784 CGTGAGCTGGGGAGTGGGCTGTGGTCGGCCCAATCGGCCCGGTGTCTACACCAATATCAG 843

Qy        841 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 900
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Db        844 CCACCACTTTGAGTGGATCCAGAAGCTGATGGCCCAGAGTGGCATGTCCCAGCCAGACCC 903

Qy        901 CTCCTGGCCGCTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 960
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Db        904 CTCCTGGCCACTACTCTTTTTCCCTCTTCTCTGGGCTCTCCCACTCCTGGGGCCGGTCTG 963

Qy        961 AGCCTACCTGAGCCCATGCAGCCTGGGGCCACTGCCAAGTCAGGCCCTGGTTCTCTTCTG 1020
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Qy       1021 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTTTTCAAAAAAAAA 1080
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Db       1024 TCTTGTTTGGTAATAAACACATTCCAGTTGATGCCTTGCAGGGCATTCTTCAAAAAAAAA 1083

Qy       1081 AAAAAAAAAAAAAAAA 1097
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Db       1084 AAAAAAAAAAAAAAAA 1100
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Job time : 683.27 secs